

Y79AA1001323//Mus musculus mRNA for GSG1, complete cds.

Y79AA1001402//Homo sapiens paraneoplastic cancer-testis-brain antigen (MA4) mRNA, partial cds.

Y79AA1001679//Homo sapiens lambda-crystallin mRNA, complete cds.

5 Y79AA1001923//Homo sapiens F-box protein Fbx22 (FBX22) gene, partial cds. Y79AA1002083//H. sapiens mRNA for MUF1 protein.

Y79AA1002307//Homo sapiens astrotactin2 (ASTN2) mRNA, complete cds.

Y79AA1002311//R. norvegicus mRNA for cytosolic resiniferatoxin-binding protein.

Y79AA1002487//Homo sapiens chromosome 5 F-box protein Fbx4 (FBX4) mRNA, complete cds.

10 **[0292]** Among the clones other than the above-mentioned, there were 36 clones that were similarly classified into the functional categories based on the results of functional domain search using the Pfam program. These clones were categorized as follows.

[0293] Clones presumably belonging to the category of secretory or membrane proteins are two clones, MAMMA1002498 and NT2RM4002287; a clone presumably belonging to the category of glycoproteins-associated proteins is a clone MAMMA1002498; clones presumably belonging to the category of signal transduction-associated proteins are 11 clones, HEMBA1001247, NT2RM2001813, NT2RM4001454, NT2RP2005140, NT2RP2005293, NT2RP3000487, NT2RP3003311, PLACE1000972, PLACE1003723, PLACE1005327, and PLACE3000124; clones presumably belonging to the category of transcription-associated proteins are 12 clones, HEMBA1003257, NT2RM2000101, NT2RM2001797, NT2RP1000101, NT2RP2002208, NT2RP3001214, NT2RP3003278, NT2RP4001235, PLACE1000050, PLACE1001716, PLACE1002499, and PLACE1007544; clones presumably belonging to the category of enzymes and/or metabolism-associated proteins are 2 clones, HEMBA1005732 and MAMMA1000402; clones presumably belonging to the category of DNA- and/or RNA-binding proteins are 4 clones, HEMBA1004596, OVARC1000148, PLACE1003334, and THYRO1001661; a clone presumably belonging to the category of protein synthesis- and/or protein transport-associated proteins is a clone, HEMBA1006284.

25 **[0294]** So far, useful information for presuming the functions is unavailable for the remaining 2511 clones. Their functions will possibly be revealed by further analyses. Names of the clones are listed below.

[0295] So far, useful information for presuming the functions is unavailable for the remaining 2511 clones. Their functions will possibly be revealed by further analyses. Names of the clones are listed below.

30 HEMBA1000042, HEMBA1000046, HEMBA1000050, HEMBA1000076, HEMBA1000193, HEMBA1000213,
HEMBA1000227, HEMBA1000231, HEMBA1000243, HEMBA1000244, HEMBA1000251, HEMBA1000264,
HEMBA1000280, HEMBA1000282, HEMBA1000288, HEMBA1000290, HEMBA1000302, HEMBA1000327,
HEMBA1000338, HEMBA1000351, HEMBA1000357, HEMBA1000376, HEMBA1000387, HEMBA1000392,
HEMBA1000396, HEMBA1000428, HEMBA1000442, HEMBA1000456, HEMBA1000459, HEMBA1000460,
HEMBA1000469, HEMBA1000497, HEMBA1000501, HEMBA1000504, HEMBA1000505, HEMBA1000508,
35 HEMBA1000519, HEMBA1000520, HEMBA1000534, HEMBA1000545, HEMBA1000557, HEMBA1000568,
HEMBA1000575, HEMBA1000594, HEMBA1000604, HEMBA1000622, HEMBA1000636, HEMBA1000655,
HEMBA1000673, HEMBA1000682, HEMBA1000686, HEMBA1000702, HEMBA1000722, HEMBA1000726,
HEMBA1000727, HEMBA1000749, HEMBA1000752, HEMBA1000769, HEMBA1000773, HEMBA1000774,
HEMBA1000843, HEMBA1000867, HEMBA1000869, HEMBA1000872, HEMBA1000876, HEMBA1000908,
40 HEMBA1000918, HEMBA1000934, HEMBA1000942, HEMBA1000943, HEMBA1000946, HEMBA1000960,
HEMBA1000968, HEMBA1000971, HEMBA1000972, HEMBA1000975, HEMBA1000985, HEMBA1000986,
HEMBA1001008, HEMBA1001009, HEMBA1001020, HEMBA1001022, HEMBA1001024, HEMBA1001026,
HEMBA1001051, HEMBA1001060, HEMBA1001080, HEMBA1001094, HEMBA1001099, HEMBA1001109,
HEMBA1001121, HEMBA1001122, HEMBA1001123, HEMBA1001133, HEMBA1001140, HEMBA1001208,
45 HEMBA1001213, HEMBA1001226, HEMBA1001235, HEMBA1001281, HEMBA1001299, HEMBA1001303,
HEMBA1001310, HEMBA1001319, HEMBA1001323, HEMBA1001326, HEMBA1001327, HEMBA1001330,
HEMBA1001361, HEMBA1001375, HEMBA1001377, HEMBA1001383, HEMBA1001388, HEMBA1001391,
HEMBA1001398, HEMBA1001411, HEMBA1001413, HEMBA1001415, HEMBA1001432, HEMBA1001433,
HEMBA1001435, HEMBA1001442, HEMBA1001450, HEMBA1001463, HEMBA1001497, HEMBA1001522,
50 HEMBA1001533, HEMBA1001566, HEMBA1001570, HEMBA1001581, HEMBA1001589, HEMBA1001608,
HEMBA1001636, HEMBA1001640, HEMBA1001647, HEMBA1001655, HEMBA1001658, HEMBA1001702,
HEMBA1001711, HEMBA1001712, HEMBA1001731, HEMBA1001745, HEMBA1001750, HEMBA1001781,
HEMBA1001784, HEMBA1001791, HEMBA1001803, HEMBA1001815, HEMBA1001820, HEMBA1001835,
HEMBA1001864, HEMBA1001888, HEMBA1001910, HEMBA1001912, HEMBA1001915, HEMBA1001918,
55 HEMBA1001939, HEMBA1001940, HEMBA1001942, HEMBA1001950, HEMBA1001964, HEMBA1001987,
HEMBA1002018, HEMBA1002022, HEMBA1002039, HEMBA1002049, HEMBA1002084, HEMBA1002100,
HEMBA1002113, HEMBA1002119, HEMBA1002160, HEMBA1002162, HEMBA1002185, HEMBA1002189,
HEMBA1002191, HEMBA1002199, HEMBA1002204, HEMBA1002229, HEMBA1002237, HEMBA1002265,

PLACE1004868//MALE STERILITY PROTEIN 2.
 PLACE1004930//Homo sapiens MDC-3.13 isoform 2 mRNA, complete cds
 PLACE1005052//Homo sapiens CGI-16 protein mRNA, complete cds.
 PLACE1005102//RING CANAL PROTEIN (KELCH PROTEIN).
 5 PLACE1005176//Homo sapiens hypothalamus protein HT001 mRNA, complete cds.
 PLACE1005187//APAG PROTEIN.
 PLACE1005331//Homo sapiens 7h3 protein mRNA, partial cds.
 PLACE1005727//Homo sapiens STRIN protein (STRIN) mRNA, complete cds.
 PLACE1006003//Homo sapiens CGI-94 protein mRNA, complete cds.
 10 PLACE1006335//Homo sapiens NY-REN-50 antigen mRNA, partial cds.
 PLACE1006385//Homo sapiens epsin 2a mRNA, complete cds.
 PLACE1006506//Homo sapiens anaphase-promoting complex subunit 4 (APC4) mRNA, complete cds.
 PLACE1007105//Homo sapiens muskelin (MKLN1) mRNA, complete cds.
 PLACE1007537//Homo sapiens ankyrin repeat-containing protein ASB-2 mRNA, complete cds.
 15 PLACE1007705//Mus musculus mRNA for Ndr1 related protein Ndr3, complete cds.
 PLACE1007791//Homo sapiens IDN3-B mRNA, complete cds.
 PLACE1007897//Homo sapiens FLASH mRNA, complete cds.
 PLACE1008080//Homo sapiens mRNA for HEXIM1 protein, complete cds.
 PLACE1008368//RING CANAL PROTEIN (KELCH PROTEIN).
 20 PLACE1008398//GENE 33 POLYPEPTIDE.
 PLACE1008465//Homo sapiens mRNA for rapa-1 (rapa gene).
 PLACE1008627//Homo sapiens mRNA for cysteine-rich protein.
 PLACE1009020//NIFS PROTEIN.
 PLACE1009060//BRO1 PROTEIN.
 25 PLACE1009186//Homo sapiens small zinc finger-like protein (TIM9b) mRNA, complete cds.
 PLACE1009443//Mus musculus F-box protein FBL8 mRNA, complete cds.
 PLACE1009571//Homo sapiens PTD002 mRNA, complete cds.
 PLACE1009670//Homo sapiens genethonin 1 mRNA, complete cds.
 PLACE1010105//RING CANAL PROTEIN (KELCH PROTEIN).
 30 PLACE1010261//SEGREGATION DISTORTER PROTEIN.
 PLACE1010310//SPIDROIN 2 (DRAGLINE SILK FIBROIN 2) (FRAGMENT).
 PLACE1010522//Homo sapiens mRNA for DEPP (decidual protein induced by progesterone), complete cds.
 PLACE1010579//Homo sapiens CED-6 protein (CED-6) mRNA, complete cds.
 PLACE1010628//Homo sapiens S164 gene, partial cds; PS1 and hypothetical protein genes, complete cds; and
 35 S171 gene, partial cds.
 PLACE1010661//TESTIS-SPECIFIC PROTEIN PBS13.
 PLACE1010761//Homo sapiens mRNA for cisplatin resistance-associated overexpressed protein, complete cds.
 PLACE1011185//INSERTION ELEMENT IS1 PROTEIN INSB.
 PLACE1011340//Homo sapiens IDN3-B mRNA, complete cds.
 40 PLACE1011586//Rattus norvegicus clone C53 CDK5 activator-binding protein mRNA, complete cds.
 PLACE2000246//RING CANAL PROTEIN (KELCH PROTEIN).
 PLACE2000411//Homo sapiens epsin 2b mRNA, complete cds.
 PLACE3000477//Homo sapiens phosphoprotein pp75 mRNA, partial cds.
 THYRO1000173//Homo sapiens AP-mu chain family member mu1B (HSMU1B) mRNA, complete cds.
 45 THYRO1000401//Human TcD37 homolog (HTcD37) mRNA, partial cds.
 THYRO1000666//Mus musculus mRNA for kinesin like protein 9.
 THYRO1001033//TRANSFORMATION-SENSITIVE PROTEIN IEF SSP 3521.
 THYRO1001347//Homo sapiens RAN binding protein 16 mRNA, complete cds.
 THYRO1001656//Homo sapiens Leman coiled-coil protein (LCCP) mRNA, complete cds.
 50 THYRO1001703//NIFR3-LIKE PROTEIN.
 THYRO1001721//RING CANAL PROTEIN (KELCH PROTEIN).
 Y79AA1000059//Homo sapiens aryl-hydrocarbon interacting protein-like 1 (AIPL1) gene, complete cds.
 Y79AA1000181//Homo sapiens CGI-01 protein mRNA, complete cds.
 Y79AA1000268//Mus musculus Nip2l mRNA, complete cds.
 55 Y79AA1000313//CALPHOTIN.
 Y79AA1000540//CELL POLARITY PROTEIN TEA1.
 Y79AA1000966//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.
 Y79AA1000985//Human centrosomal protein kendrin mRNA, complete cds.

- NT2RP3001712//Homo sapiens HP1-BP74 protein mRNA, complete cds.
 NT2RP3001819//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3001854//Homo sapiens novel retinal pigment epithelial cell protein (NORPEG) mRNA, complete cds.
 NT2RP3001931//Rattus norvegicus clone C48 CDK5 activator-binding protein mRNA, complete cds.
 5 NT2RP3002273//SCD6 PROTEIN.
 NT2RP3002631//Homo sapiens Ran binding protein 11 mRNA, complete cds.
 NT2RP3002682//Homo sapiens CGI-145 protein mRNA, complete cds.
 NT2RP3002770//MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116.
 NT2RP3002818//INSERTION ELEMENT IS2A HYPOTHETICAL 48.2 KD PROTEIN.
 10 NT2RP3002948//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3002972//Halocynthia roretzi mRNA for HrPET-1, complete cds.
 NT2RP3003032//Homo sapiens okadaic acid-inducible and cAMP-regulated phosphoprotein 19 (ARPP-19) mRNA, complete cds.
 NT2RP3003290//Mus musculus mRNA for Ndr1 related protein Ndr3, complete cds.
 15 NT2RP3003411//Mus musculus COP9 complex subunit 7b (COPS7b) mRNA, complete cds.
 NT2RP3003491//Drosophila melanogaster Pelle associated protein Pellino (Pli) mRNA, complete cds.
 NT2RP3003500//SCY1 PROTEIN.
 NT2RP3003726//Homo sapiens spermatogenesis associated PD1 mRNA, complete cds.
 NT2RP3004348//R. norvegicus mRNA for cytosolic resiniferatoxin-binding protein.
 20 NT2RP3004507//MOB1 PROTEIN (MPS1 BINDER 1).
 NT2RP4000129//Xenopus laevis F-box protein 28 (Fbx28) mRNA, partial cds.
 NT2RP4000498//MOB1 PROTEIN (MPS1 BINDER 1).
 NT2RP4000528//NPL4 PROTEIN.
 NT2RP4000737//Mus musculus F-box protein FBL10 mRNA, partial cds.
 25 NT2RP4000979//Homo sapiens putative HIV-1 infection related protein mRNA, partial cds.
 NT2RP4001010//Rattus norvegicus PSD-95/SAP90-associated protein-4 mRNA, complete cds.
 NT2RP4001207//Homo sapiens Ran binding protein 11 mRNA, complete cds.
 NT2RP4001228//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP4001260//Homo sapiens F-box protein Fbx21 (FBX21) mRNA, complete cds.
 30 NT2RP4001339//Homo sapiens mRNA for AMMERC1 protein.
 NT2RP4001351//Human ovarian cancer downregulated myosin heavy chain homolog (Doc1) mRNA, complete cds.
 NT2RP4001474//Xenopus laevis putative Zic3 binding protein mRNA, complete cds.
 NT2RP4001966//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
 35 NT2RP4002018//RING CANAL PROTEIN (KELCH PROTEIN).
 OVARC1000209//Oryza sativa submergence induced protein 2A mRNA, complete cds.
 OVARC1000876//MOB1 PROTEIN (MPS1 BINDER 1).
 OVARC1001065//Homo sapiens CGI-12 protein mRNA, complete cds.
 OVARC1001092//Homo sapiens mRNA for JM5 protein, complete CDS (clone IMAGE 53337, LLNLc110F185707 (RZPD Berlin) and LLNLc110G0913Q7 (RZPD Berlin)).
 40 OVARC1001419//Homo sapiens GOK (STIM1) mRNA, complete cds.
 OVARC1001555//NGG1-INTERACTING FACTOR 3.
 OVARC1001711//CORNIFIN B (SMALL PROLINE-RICH PROTEIN 1B) (SPR1B) (SPR1 B).
 OVARG1001943//Mus musculus DEBT-91 mRNA, complete cds.
 45 PLACE1000004//Homo sapiens IDN3-B mRNA, complete cds.
 PLACE1000066//SSU72 PROTEIN.
 PLACE1000610//MSN5 PROTEIN.
 PLACE1000636//MALE STERILITY PROTEIN 2.
 PLACE1000769//Homo sapiens CGI-18 protein mRNA, complete cds.
 50 PLACE1000987//Rattus norvegicus late gestation lung 2 protein (Lgl2) mRNA, complete cds.
 PLACE1001036//Homo sapiens mRNA for alpha integrin binding protein 63, partial.
 PLACE1001845//Mus musculus cyclin ania-6a mRNA, complete cds.
 PLACE1001920//Homo sapiens MDC-3.13 isoform 2 mRNA, complete cds.
 PLACE1002665//Mus musculus enhancer of polycomb (Epc1) mRNA, complete cds.
 55 PLACE1003602//Homo sapiens mRNA expressed in placenta.
 PLACE1003611//Homo sapiens anaphase-promoting complex subunit 4 (APC4) mRNA, complete cds.
 PLACE1004256//Mus musculus short coiled coil protein SCOCO (Scoc) mRNA, complete cds.
 PLACE1004550//Homo sapiens CGI-20 protein mRNA, complete cds.

- NT2RM4001731//Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds.
 NT2RM4001969//R.norvegicus mRNA for IP63 protein.
 NT2RM4002034//Homo sapiens hiwi mRNA, partial cds.
 NT2RM4002075//RING CANAL PROTEIN (KELCH PROTEIN).
 5 NT2RM4002226//GTPASE ACTIVATING PROTEIN ROTUND.
 NT2RP1000040//Mus musculus donson protein (Donson) mRNA, partial cds.
 NT2RP1000363//R.norvegicus LL5 mRNA.
 NT2RP1000481//Homo sapiens antigen NY-CO-3 (NY-CO-3) mRNA, partial cds.
 NT2RP1000513//Human NifU-like protein (hNifU) mRNA, partial cds.
 10 NT2RP1000733//Human mRNA for GSPT1-TK protein, complete cds.
 NT2RP1000860//Homo sapiens KLO4P mRNA, complete cds.
 NT2RP1000954//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP1001011//Drosophila melanogaster putative 43 kDa protein (TH1) mRNA, complete cds.
 NT2RP1001395//Homo sapiens COP9 complex subunit 7a mRNA, complete cds.
 15 NT2RP1001457//Homo sapiens partial mRNA for beta-transducin family protein (putative).
 NT2RP1001494//MALE STERILITY PROTEIN 2.
 NT2RP2000054//Homo sapiens putative ring zinc finger protein NY-REN-43 antigen mRNA, complete cds.
 NT2RP2000067//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
 NT2RP2000133//Homo sapiens Leman coiled-coil protein (LCCP) mRNA, complete cds.
 20 NT2RP2000157//MLO2 PROTEIN.
 NT2RP2000764//NIFS PROTEIN.
 NT2RP2000965//Homo sapiens mRNA for fls353, complete cds.
 NT2RP2001839//SCY1 PROTEIN.
 NT2RP2001883//Homo sapiens CGI-01 protein mRNA, complete cds.
 25 NT2RP2001976//Mus musculus calmodulin-binding protein SHA1 (Sha1) mRNA, complete cds.
 NT2RP2001985//Homo sapiens high-risk human papilloma viruses E6 oncoproteins targeted protein E6TP1 alpha mRNA, complete cds.
 NT2RP2002185//Homo sapiens ubiquilin mRNA, complete cds.
 NT2RP2002442//HESA PROTEIN.
 30 NT2RP2002727//Rattus norvegicus tulip 2 mRNA, complete cds.
 NT2RP2002741//Homo sapiens mRNA for Neuroblastoma, complete cds.
 NT2RP2002986//Homo sapiens mRNA for Kelch motif containing protein, complete cds.
 NT2RP2003121//Mus musculus enhancer of polycomb (Epc1) mRNA, complete cds.
 NT2RP2003265//Homo sapiens CGI-53 protein mRNA, complete cds.
 35 NT2RP2003272//Homo sapiens ubiquilin mRNA, complete cds.
 NT2RP2003857//MYOTROPHIN (V-1 PROTEIN) (GRANULE CELL DIFFERENTIATION PROTEIN).
 NT2RP2003871//Homo sapiens transposon-derived Buster1 transposase-like protein gene, complete cds.
 NT2RP2004425//Mus musculus axotrophin mRNA, complete cds.
 NT2RP2004476//Homo sapiens cyclin L ania-6a mRNA, complete cds.
 40 NT2RP2004710//Mus musculus formin binding protein 30 mRNA, complete cds.
 NT2RP2004816//H58 PROTEIN.
 NT2RP2005441//Homo sapiens hypothalamus protein HT002 mRNA, complete cds.
 NT2RP2005490//Mus musculus D3Mm3e (D3Mm3e) mRNA, complete cds.
 NT2RP2005620//Homo sapiens epsin 2a mRNA, complete cds.
 45 NT2RP2005654//CYSTEINE STRING PROTEIN (CCCS1).
 NT2RP2005675//Homo sapiens growth suppressor related (DOC-1R) mRNA, complete cds.
 NT2RP2005753//Homo sapiens I-1 receptor candidate protein mRNA, complete cds.
 NT2RP2005841//Homo sapiens mRNA for ALEX3, complete cds.
 NT2RP2006598//Homo sapiens retinoid x receptor interacting protein mRNA, complete cds.
 50 NT2RP3000047//NPL4 PROTEIN.
 NT2RP3000233//RING CANAL PROTEIN (KELCH PROTEIN).
 NT2RP3000868//Human ovarian cancer downregulated myosin heavy chain homolog (Doc1) mRNA, complete cds.
 NT2RP3000869//Drosophila melanogaster AAA family protein Bor (bor) mRNA, complete cds.
 55 NT2RP3001399//SSU72 PROTEIN.
 NT2RP3001407//SCY1 PROTEIN.
 NT2RP3001457//Drosophila melanogaster Melted (melt) mRNA, partial cds.
 NT2RP3001587//Human anthracycline-associated resistance ARX mRNA, complete cds.

- HEMBA1006398//Human L1 element L1.6 putative p150 gene, complete cds.
HEMBA1006445//Homo sapiens putative tumor supressor NOEY2 mRNA, complete cds.
HEMBA1007174//Homo sapiens epsin 2b mRNA, complete cds.
5 HEMBA1007251//Homo sapiens F-box protein FBX29 (FBX29) mRNA, partial cds. HEMBB1000036//Homo sapiens CGI-51 protein mRNA, complete cds.
HEMBB1000144//GUANYLATE CYCLASE ACTIVATING PROTEIN 2 (GCAP 2) (RETINAL GUANYLYL CYCLASE ACTIVATOR PROTEIN P24).
HEMBB1000973//Mus musculus schlafen3 (Sln3) mRNA, complete cds.
HEMBB1001058//Homo sapiens neuronal thread protein AD7c-NTP mRNA, complete cds
10 HEMBB1001234//65 KD YES-ASSOCIATED PROTEIN (YAP65).
HEMBB1001288//COPPER HOMEOSTASIS PROTEIN CUTC.
HEMBB1001331//Mus musculus mRNA for hepatoma-derived growth factor, complete cds, strain:BALB/c.
HEMBB1001384//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.
HEMBB1002266//NEURONAL PROTEIN.
15 HEMBB1002510//GYP7 PROTEIN.
HEMBB1002705//Homo sapiens CGI-27 protein mRNA, complete cds.
MAMMA1000055//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TEST)].
MAMMA1000625//GYP7 PROTEIN.
MAMMA1001075//Homo sapiens CGI-72 protein mRNA, complete cds.
20 MAMMA1001181//ABC1 PROTEIN HOMOLOG PRECURSOR.
MAMMA1001259//Mus musculus F-box protein FBX18 mRNA, partial cds.
MAMMA1001730//Homo sapiens brain and nasopharyngeal carcinoma susceptibility protein NSG-x mRNA, partial cds.
MAMMA1002143//Homo sapiens Cdc42 effector protein 4 mRNA, complete cds.
25 MAMMA1002699//Rattus norvegicus EH domain binding protein Epsin mRNA, complete cds.
MAMMA1002972//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS27.
MAMMA1003113//Mus musculus COP9 complex subunit 7a (COPS7a) mRNA, complete cds.
NT2RM1000118//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT).
30 NT2RM1000186//CALCINEURIN B SUBUNIT (PROTEIN PHOSPHATASE 2B REGULATORY SUBUNIT) (CALCINEURIN REGULATORY SUBUNIT).
NT2RM1000244//Homo sapiens TRAF4 associated factor 1 mRNA, partial cds.
NT2RM1000421//RIBONUCLEASE INHIBITOR.
NT2RM1000499//Caenorhabditis elegans mRNA for centaurin gamma 1A.
35 NT2RM1000623//RIBONUCLEASE INHIBITOR.
NT2RM1000883//Homo sapiens I-1 receptor candidate protein mRNA, complete cds.
NT2RM2000502//Rattus norvegicus W307 mRNA, complete cds.
NT2RM2000599//Homo sapiens F-box protein Lilina (LILINA) mRNA, complete cds.
NT2RM2000718//Homo sapiens endocrine regulator mRNA, complete cds.
40 NT2RM2001065//Homo sapiens COP9 complex subunit 4 mRNA, complete cds.
NT2RM2001196//PROLINE-RICH PROTEIN MP-3 (FRAGMENT).
NT2RM2001983//Homo sapiens RGS-GAIP interacting protein GIPC mRNA, complete cds.
NT2RM2002109//Homo sapiens glioma amplified on chromosome 1 protein (GAC1) mRNA, complete cds.
NT2RM2002142//GASTRULATION SPECIFIC PROTEIN G12.
45 NT2RM4000030//LAS1 PROTEIN.
NT2RM4000139//R.norvegicus trg mRNA.
NT2RM4000156//H. sapiens HPBRII-7 gene.
NT2RM4000386//Mus musculus ODZ3 (Odz3) mRNA, partial cds.
NT2RM4000590//RING CANAL PROTEIN (KELCH PROTEIN).
50 NT2RM4001047//MO25 PROTEIN.
NT2RM4001155//ADRENAL MEDULLA 50 KD PROTEIN.
NT2RM4001256//Xenopus laevis putative Zic3 binding protein mRNA, complete cds.
NT2RM4001320//Homo sapiens mRNA for Neuroblastoma, complete cds.
NT2RM4001340//UTR4 PROTEIN (UNKNOWN TRANSCRIPT 4 PROTEIN).
55 NT2RM4001347//Homo sapiens NY-REN-25 antigen mRNA, partial cds.
NT2RM4001371//Homo sapiens IDN3 mRNA, partial cds.
NT2RM4001582//Mus musculus COP9 complex subunit 7b (COPS7b) mRNA, complete cds.
NT2RM4001611//SIS2 PROTEIN (HALOTOLERANCE PROTEIN HAL3).

NT2RM2001896, NT2RM2002145, NT2RM4000215, NT2RM4001714, NT2RP1000163, NT2RP1000333,
NT2RP1000439, NT2RP2000346, NT2RP2001397, NT2RP2002595, NT2RP2003177, NT2RP2003596,
NT2RP2003912, NT2RP2004396, NT2RP2005037, NT2RP2005520, NT2RP2005669, NT2RP2005835,
NT2RP3001730, NT2RP3002081, NT2RP4000210, NT2RP4000415, NT2RP4001414, NT2RP4001634,
5 OVARC1000013, OVARC1000937, PLACE1001383, PLACE1002433, PLACE1004316, PLACE1005287,
PLACE1008808, PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394,

[0289] The following 36 clones presumably belong to the category of embryogenesis- and/or development-associated proteins.

HEMBA1000518, HEMBA1001847, HEMBA1001869, HEMBA1003545, HEMBA1004973, HEMBB1002442,
10 MAMMA1001837, NT2RM2001670, NT2RM4000046, NT2RM4000531, NT2RM4001140, NT2RM4001858,
NT2RP2002078, NT2RP2004187, NT2RP2006436, NT2RP3000603, NT2RP3000994, NT2RP3001580,
NT2RP3001708, NT2RP3003071, NT2RP3004472, NT2RP3004617, NT2RP4000246, NT2RP4001567,
OVARC1000304, OVARC1000746, PLACE1000793, PLACE1002532, PLACE1003258, PLACE1003625,
PLACE1004460, PLACE1009622, PLACE4000558, THYRO1000085, Y79AA1001391, Y79AA1001692,

15 [0290] The following 30 clones presumably belong to cellular defense-associated proteins.

HEMBA1000005, HEMBA1000531, HEMBA1003417, HEMBA1006253, NT2RM4000354, NT2RM4001880,
NT2RP1000333, NT2RP1000493, NT2RP2000006, NT2RP2000045, NT2RP2000809, NT2RP2001536,
NT2RP2002464, NT2RP2004920, NT2RP2005037, NT2RP3000590, NT2RP3001426, NT2RP3002062,
NT2RP3002785, NT2RP3004262, NT2RP4001555, NT2RP4001638, PLACE1006958, PLACE1008275,
20 PLACE1009113, PLACE1011858, PLACE4000014, THYRO1000684, Y79AA1002139, Y79AA1002229,

[0291] Although it is unclear whether or not 261 clones out of clones other than the above-mentioned clones belong to any of the above-described categories, these clones are predicted to have some functions, based on the homology search using the full-length sequences thereof. The clone names and the gene definitions found in the result of homology search are shown below, separated with a double-slash mark, //.

25 HEMBA1000030//Homo sapiens ARF GTPase-activating protein GIT1 mRNA, complete cds.

HEMBA1000307//CARNITINE DEFICIENCY-ASSOCIATED PROTEIN EXPRESSED IN VENTRICLE 1

30 HEMBA1000333//Homo sapiens F-box protein Fbx21 (FBX21) mRNA, complete cds.

HEMBA1000488//RING CANAL PROTEIN (KELCH PROTEIN).

HEMBA1000523//TESTIS-SPECIFIC PROTEIN PBS13.

HEMBA1001197//Homo sapiens rap2 interacting protein x mRNA, complete cds.

HEMBA1001302//Homo sapiens calcium binding protein precursor, mRNA, complete cds.

35 HEMBA1001455//Mus musculus transposon-derived Buster2 transposase-like protein gene, partial cds.

HEMBA1001675//VACUOLAR PROTEIN SORTING-ASSOCIATED PROTEIN VPS9.

HEMBA1001714//Homo sapiens mRNA for ATPase inhibitor precursor, complete cds.

HEMBA1001744//SCY1 PROTEIN.

HEMBA1001967//Homo sapiens NY-REN-57 antigen mRNA, partial cds.

40 HEMBA1002151//Rattus norvegicus p34 mRNA, complete cds.

HEMBA1002215//TESTIN 2 (TES2) [CONTAINS: TESTIN 1 (TES1)].

HEMBA1002458//OVARIAN GRANULOSA CELL 13.0 KD PROTEIN HGR74.

HEMBA1002777//Fugu rubripes BAW (BAW) mRNA, complete cds.

HEMBA1003098//Homo sapiens NY-REN-6 antigen mRNA, partial cds.

45 HEMBA1003199//Homo sapiens chromosome 5 F-box protein Fbx4 (FBX4) mRNA, complete cds.

HEMBA1003615//Homo sapiens ART-4 mRNA, complete cds.

HEMBA1003836//MOB1 PROTEIN (MPS1 BINDER 1).

HEMBA1004295//Homo sapiens NY-REN-25 antigen mRNA, partial cds.

HEMBA1004573//Homo sapiens mRNA for HELG protein.

50 HEMBA1004604//Homo sapiens COP9 complex subunit 7a mRNA, complete cds.

HEMBA1004795//CDC4-LIKE PROTEIN (FRAGMENT).

HEMBA1005101//Homo sapiens SYT interacting protein SIP mRNA, complete cds.

HEMBA1005201//Homo sapiens CGI-07 protein mRNA, complete cds.

HEMBA1005206//Drosophila simulans anon73B1 gene and Su(P) gene.

55 HEMBA1005530//Homo sapiens anaphase-promoting complex subunit 7 (APC7) mRNA, complete cds.

HEMBA1005666//Homo sapiens mRNA for DIPB protein.

HEMBA1005990//Homo sapiens I-1 receptor candidate protein mRNA, complete cds.

HEMBA1006268//Homo sapiens HQOQ24c mRNA, complete cds.

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	HEMBA1004276,	HEMBA1004734,	HEMBA1004847,	HEMBA1004929,	HEMBA1004930,	HEMBA1005047,
	HEMBA1005202,	HEMBA1006031,	HEMBA1006272,	HEMBA1006474,	HEMBA1006652,	HEMBA1006914,
	HEMBA1006973,	HEMBA1007224,	HEMBA1000915,	HEMBA1001112,	HEMBA1001137,	HEMBA1001736,
	HEMBA1001831,	HEMBA1001915,	MAMMA1000085,	MAMMA1000734,	MAMMA1001008,	MAMMA1002170,
5	MAMMA1002219,	MAMMA1002236,	MAMMA1002619,	NT2RM1000661,	NT2RM1000833,	NT2RM2000092,
	NT2RM2000504,	NT2RM2000577,	NT2RM2000821,	NT2RM2001201,	NT2RM2001592,	NT2RM2001613,
	NT2RM2001648,	NT2RM2001730,	NT2RM2001760,	NT2RM2002055,	NT2RM4000155,	NT2RM4000169,
	NT2RM4000344,	NT2RM4000356,	NT2RM4000421,	NT2RM4000712,	NT2RM4001054,	NT2RM4001203,
	NT2RM4001382,	NT2RM4001444,	NT2RM4002062,	NT2RM4002205,	NT2RM4002623,	NT2RP1000326,
10	NT2RP1000522,	NT2RP1000547,	NT2RP1000746,	NT2RP1000947,	NT2RP1001569,	NT2RP2000147,
	NT2RP2000710,	NT2RP2000880,	NT2RP2000943,	NT2RP2001290,	NT2RP2001392,	NT2RP2001601,
	NT2RP2001613,	NT2RP2001660,	NT2RP2001740,	NT2RP2002124,	NT2RP2002606,	NT2RP2002862,
	NT2RP2002959,	NT2RP2002980,	NT2RP2003137,	NT2RP2003158,	NT2RP2003391,	NT2RP2003394,
	NT2RP2003401,	NT2RP2003433,	NT2RP2003704,	NT2RP2003713,	NT2RP2003737,	NT2RP2003760,
15	NT2RP2003981,	NT2RP2004366,	NT2RP2004389,	NT2RP2004791,	NT2RP2005012,	NT2RP2005116,
	NT2RP2005360,	NT2RP2005763,	NT2RP2005784,	NT2RP3000366,		
	NT2RP3000759,	NT2RP3000968,	NT2RP3001113,	NT2RP3001690,	NT2RP3002045,	NT2RP3002151,
	NT2RP3002529,	NT2RP3002671,	NT2RP3003301,	NT2RP3003846,	NT2RP3003876,	NT2RP3004209,
	NT2RP4000370,	NT2RP4000457,	NT2RP4000879,	NT2RP4000927,	NT2RP4001041,	NT2RP4001117,
20	NT2RP4001313,	NT2RP4001315,	NT2RP4001574,	NT2RP4001592,	OVARC1000013,	OVARC1000071,
	OVARC1000085,	OVARC1000465,	OVARC1000564,	OVARC1000771,	OVARC1000862,	OVARC1001171,
	OVARC1001180,	OVARC1001342,	PLACE1000007,	PLACE1000061,	PLACE1000081,	PLACE1000492,
	PLACE1000863,	PLACE1001092,	PLACE1001748,	PLACE1002090,	PLACE1003174,	PLACE1003915,
	PLACE1004104,	PLACE1004270,	PLACE1004743,	PLACE1005557,	PLACE1005813,	PLACE1006170,
25	PLACE1006488,	PLACE1006829,	PLACE1007706,	PLACE1007729,	PLACE1008273,	PLACE1008402,
	PLACE1008790,	PLACE1008813,	PLACE1009094,	PLACE1009130,	PLACE1009477,	PLACE1009721,
	PLACE1009845,	PLACE1010074,	PLACE1010547,	PLACE1011109,	PLACE1011229,	PLACE1011477,
	PLACE1012031,	PLACE2000404,	PLACE3000059,	PLACE3000121,	PLACE4000269,	PLACE4000654,
	SKNMC1000011,	THYRO1000983,	THYRO1001003,	THYRO1001313,	Y79AA1000560,	Y79AA1000784,
30	Y79AA1000968,	Y79AA1001493,	Y79AA1001875,	Y79AA1002027,	Y79AA1002209,	
	[0287] The following 130 clones presumably belong to cytoskeletal-associated proteins.					
	HEMBA1000156,	HEMBA1000168,	HEMBA1000411,	HEMBA1000588,	HEMBA1001043,	HEMBA1001651,
	HEMBA1001661,	HEMBA1002102,	HEMBA1002161,	HEMBA1002939,	HEMBA1003235,	HEMBA1003581,
	HEMBA1004499,	HEMBA1004534,	HEMBA1004697,	HEMBA1004929,	HEMBA1004972,	HEMBA1005582,
35	HEMBA1005595,	HEMBA1006344,	HEMBA1006737,	HEMBA1001175,	HEMBA1001282,	HEMBA1001562,
	HEMBA1001802,	MAMMA1000824,	MAMMA1001041,	MAMMA1001576,	MAMMA1001679,	MAMMA1001735,
	MAMMA1002297,	MAMMA1002351,	MAMMA1002622,	MAMMA1002637,	MAMMA1003127,	NT2RM1000850,
	NT2RM1000898,	NT2RM2000030,	NT2RM2000260,	NT2RM2000691,	NT2RM2001324,	NT2RM4000169,
	NT2RM4000229,	NT2RM4000515,	NT2RM4001217,	NT2RP1000202,	NT2RP1000348,	NT2RP1000460,
40	NT2RP1000478,	NT2RP1001033,	NT2RP1001294,	NT2RP1001302,	NT2RP2000070,	NT2RP2000812,
	NT2RP2000814,	NT2RP2001168,	NT2RP2001245,	NT2RP2001634,	NT2RP2001900,	NT2RP2003307,
	NT2RP2003394,	NT2RP2004041,	NT2RP2004242,	NT2RP2004538,	NT2RP2004587,	NT2RP2004681,
	NT2RP2004732,	NT2RP2004978,	NT2RP2005491,	NT2RP2005531,	NT2RP2005712,	NT2RP2006275,
	NT2RP3000753,	NT2RP3001113,	NT2RP3001216,	NT2RP3001239,	NT2RP3001272,	NT2RP3001554,
45	NT2RP3001690,	NT2RP3001799,	NT2RP3002688,	NT2RP3003061,	NT2RP3003185,	NT2RP3003230,
	NT2RP3004569,	NT2RP3004578,	NT2RP4001004,	NT2RP4001086,	NT2RP4001256,	NT2RP4001567,
	NT2RP4001927,	OVARC1000001,	OVARC1000106,	OVARC1000437,	OVARC1000520,	OVARC1000679,
	OVARC1001731,	OVARC1002050,	PLACE1001104,	PLACE1002571,		
	PLACE1002591,	PLACE1002655,	PLACE1002714,	PLACE1003625,	PLACE1005287,	PLACE1006552,
50	PLACE1007946,	PLACE1008426,	PLACE1010148,	PLACE1010547,	PLACE1010743,	PLACE1010896,
	PLACE1010960,	PLACE1011310,	PLACE1011922,	PLACE2000216,	PLACE2000274,	PLACE2000371,
	PLACE2000458,	PLACE3000145,	PLACE3000416,	PLACE4000009,	THYRO1000132,	THYRO1001405,
	THYRO1001458,	Y79AA1000368,	Y79AA1000794,	Y79AA1000833,	Y79AA1000962,	Y79AA1002208,
	[0288] The following 54 clones presumably belong to cell division-associated and/or cell proliferation-associated proteins.					
55	HEMBA1001019,	HEMBA1001595,	HEMBA1002363,	HEMBA1002997,	HEMBA1003136,	HEMBA1003369,
	HEMBA1004131,	HEMBA1004354,	HEMBA1005621,	HEMBA1000037,	HEMBA1000264,	MAMMA1001768,
	MAMMA1002769,	NT2RM1000354,	NT2RM1000430,	NT2RM1000874,	NT2RM2001256,	NT2RM2001743,

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	HEMBA1006248,	HEMBA1006278,	HEMBA1006283,	HEMBA1006359,	HEMBA1006652,	HEMBA1007087,
	HEMBA1007194,	HEMBA1000264,	HEMBA1000789,	HEMBA1001011,	HEMBA1001482,	HEMBA1001736,
	HEMBA1001749,	HEMBA1001839,	HEMBA1002217,	MAMMA1000183,	MAMMA1000284,	MAMMA1000731,
	MAMMA1001105,	MAMMA1001222,	MAMMA1001260,	MAMMA1001743,	MAMMA1001837,	MAMMA1002385,
5	MAMMA1002617,	MAMMA1002869,	MAMMA1002937,	MAMMA1003011,	NT2RM1000086,	NT2RM1000539,
	NT2RM1000555,	NT2RM1000666,	NT2RM1000691,	NT2RM1000826,	NT2RM1000885,	NT2RM1001059,
	NT2RM1001092,	NT2RM2000371,	NT2RM2000624,	NT2RM2000735,	NT2RM2001105,	NT2RM2001424,
	NT2RM2001575,	NT2RM2001605,	NT2RM2001670,	NT2RM2001771,	NT2RM2001823,	NT2RM2001989,
	NT2RM2002004,	NT2RM2002014,	NT2RM2002088,	NT2RM2002091,	NT2RM4000046,	NT2RM4000104,
10	NT2RM4000167,	NT2RM4000191,	NT2RM4000202,	NT2RM4000531,	NT2RM4000595,	NT2RM4000733,
	NT2RM4000751,	NT2RM4000996,	NT2RM4001092,	NT2RM4001140,		
	NT2RM4001178,	NT2RM4001200,	NT2RM4001483,	NT2RM4001592,	NT2RM4001783,	NT2RM4001823,
	NT2RM4001828,	NT2RM4001858,	NT2RM4001880,	NT2RM4001979,	NT2RM4002093,	NT2RM4002109,
	NT2RP1000470,	NT2RP1000493,	NT2RP1000574,	NT2RP1000902,	NT2RP1000966,	NT2RP1001013,
15	NT2RP1001073,	NT2RP1001080,	NT2RP2000008,	NT2RP2000153,	NT2RP2000258,	NT2RP2000297,
	NT2RP2001127,	NT2RP2001174,	NT2RP2001233,	NT2RP2001511,	NT2RP2001756,	NT2RP2001869,
	NT2RP2002079,	NT2RP2002099,	NT2RP2002503,	NT2RP2002591,	NT2RP2002939,	NT2RP2003157,
	NT2RP2003329,	NT2RP2003347,	NT2RP2003480,	NT2RP2003522,	NT2RP2003564,	NT2RP2003714,
	NT2RP2004187,	NT2RP2004568,	NT2RP2004920,	NT2RP2005003,	NT2RP2005139,	NT2RP2005168,
20	NT2RP2005436,	NT2RP2005496,	NT2RP2005701,	NT2RP2005763,	NT2RP2005776,	NT2RP2005942,
	NT2RP2006043,	NT2RP2006436,	NT2RP2006464,	NT2RP3000050,	NT2RP3000512,	NT2RP3000527,
	NT2RP3000562,	NT2RP3000590,	NT2RP3000603,	NT2RP3000624,	NT2RP3000632,	NT2RP3000994,
	NT2RP3001057,	NT2RP3001107,	NT2RP3001120,	NT2RP3001150,	NT2RP3001155,	NT2RP3001338,
	NT2RP3001398,	NT2RP3001472,	NT2RP3001672,	NT2RP3001688,	NT2RP3001724,	NT2RP3001792,
25	NT2RP3001855,	NT2RP3002165,	NT2RP3002399,	NT2RP3002876,	NT2RP3003138,	NT2RP3003193,
	NT2RP3003251,	NT2RP3003327,	NT2RP3003555,	NT2RP3004013,	NT2RP3004078,	NT2RP3004428,
	NT2RP3004490,	NT2RP3004566,	NT2RP3004594,	NT2RP3004617,	NT2RP3004618,	NT2RP4000111,
	NT2RP4000398,	NT2RP4000455,	NT2RP4000518,	NT2RP4000648,	NT2RP4000865,	NT2RP4000929,
	NT2RP4001080,	NT2RP4001095,	NT2RP4001213,	NT2RP4001433,	NT2RP4001568,	NT2RP4001696,
30	NT2RP4001753,	NT2RP4001838,	NT2RP4001938,	NT2RP4002078,	OVARC1000006,	OVARC1000087,
	OVARC1000241,	OVARC1000746,	OVARC1000846,	OVARC1001232,	OVARC1001271,	OVARC1001306,
	OVARC1001987,	OVARC1002112,	PLACE1000406,	PLACE1000583,	PLACE1000979,	PLACE1001118,
	PLACE1001632,	PLACE1001739,	PLACE1002438,	PLACE1002532,	PLACE1002775,	PLACE1002834,
	PLACE1003302,	PLACE1003519,	PLACE1003605,	PLACE1003704,	PLACE1003738,	PLACE1003885,
35	PLACE1004471,	PLACE1004564,	PLACE1004814,	PLACE1005584,	PLACE1005876,	PLACE1005951,
	PLACE1006196,	PLACE1006482,	PLACE1006488,	PLACE1006531,	PLACE1006917,	PLACE1007346,
	PLACE1007547,	PLACE1007598,	PLACE1007688,	PLACE1007969,	PLACE1008132,	PLACE1009099,
	PLACE1009246,	PLACE1009398,	PLACE1009476,	PLACE1009622,	PLACE1010053,	PLACE1010194,
	PLACE1010702,	PLACE1010870,	PLACE1011056,	PLACE1011114,	PLACE1011433,	PLACE2000427,
40	PLACE3000009,	PLACE3000169,	PLACE4000014,	PLACE4000156,	PLACE4000192,	PLACE4000261,
	PLACE4000489,	SKNMC1000091,	THYRO1000085,	THYRO1000242,	THYRO1000501,	THYRO1001100,
	THYRO1001189,	THYRO1001809,	Y79AA1000037,	Y79AA1000349,	Y79AA1000752,	Y79AA1001211,
	Y79AA1001312,	Y79AA1001391,	Y79AA1001613,	Y79AA1002103,	Y79AA1002472,	Y79AA1002482,
	[0285] The following 66 clones presumably belong to the category of RNA synthesis-associated proteins.					
45	HEMBA1000591,	HEMBA1001579,	HEMBA1003179,	HEMBA1003591,	HEMBA1006278,	HEMBA1000226,
	NT2RM1000187,	NT2RM1000852,	NT2RM2000624,	NT2RM2001989,	NT2RM2002100,	NT2RM4000191,
	NT2RM4001178,	NT2RM4002093,	NT2RP1000035,	NT2RP1000272,	NT2RP1000470,	NT2RP1001080,
	NT2RP2000153,	NT2RP2002928,	NT2RP2003157,	NT2RP2004568,	NT2RP2005126,	NT2RP2005436,
	NT2RP2005539,	NT2RP2005605,	NT2RP2005776,	NT2RP2005942,	NT2RP2006043,	NT2RP2006238,
50	NT2RP3000361,	NT2RP3000397,	NT2RP3001671,	NT2RP3004504,	NT2RP4000078,	NT2RP4000111,
	NT2RP4000481,	NT2RP4000518,	NT2RP4000614,	NT2RP4000929,	NT2RP4001696,	NT2RP4002058,
	OVARC1001232,	OVARC1001577,	PLACE1000406,	PLACE1000596,	PLACE1000755,	PLACE1001739,
	PLACE1003704,	PLACE1003885,	PLACE1004564,	PLACE1004814,	PLACE1004902,	PLACE1005373,
	PLACE1005646,	PLACE1005876,	PLACE1006196,	PLACE1006626,	PLACE1006878,	PLACE1006917,
55	PLACE1009476,	PLACE1009925,	PLACE1010194,	PLACE1011114,	THYRO1000121,	Y79AA1001963,
	[0286] The following 183 clones presumably belong to protein synthesis-associated and/or protein transport-associated proteins.					
	HEMBA1000012,	HEMBA1000141,	HEMBA1000592,	HEMBA1003617,	HEMBA1003773,	HEMBA1004202,

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	HEMBA1002569,	HEMBA1002935,	HEMBA1002951,	HEMBA1002999,	HEMBA1003408,	HEMBA1003545,
	HEMBA1003662,	HEMBA1003684,	HEMBA1003690,	HEMBA1003760,	HEMBA1004203,	HEMBA1004321,
	HEMBA1004353,	HEMBA1004479,	HEMBA1004973,	HEMBA1005219,	HEMBA1005359,	HEMBA1005558,
	HEMBA1005931,	HEMBA1006278,	HEMBA1006283,	HEMBA1006359,	HEMBA1006485,	HEMBA1007087,
5	HEMBB1000226,	HEMBB1000789,	HEMBB1001011,	HEMBB1001056,	HEMBB1001242,	HEMBB1001482,
	HEMBB1001915,	HEMBB1002134,	HEMBB1002217,	MAMMA1000183,	MAMMA1000731,	MAMMA1001105,
	MAMMA1001222,	MAMMA1001260,	MAMMA1001633,	MAMMA1001743,	MAMMA1001837,	MAMMA1002617,
	MAMMA1002869,	MAMMA1002937,	MAMMA1003011,	NT2RM1000086,	NT2RM1000187,	NT2RM1000666,
	NT2RM1000885,	NT2RM1000894,	NT2RM1001059,	NT2RM1001092,	NT2RM2000013,	NT2RM2000588,
10	NT2RM2000624,	NT2RM2000735,	NT2RM2000740,	NT2RM2001105,	NT2RM2001635,	NT2RM2001670,
	NT2RM2001771,	NT2RM2001823,	NT2RM2001936,	NT2RM2001989,	NT2RM2002004,	NT2RM2002088,
	NT2RM2002091,	NT2RM4000024,	NT2RM4000046,	NT2RM4000104,	NT2RM4000202,	NT2RM4000215,
	NT2RM4000290,	NT2RM4000531,	NT2RM4000751,	NT2RM4000996,	NT2RM4001092,	NT2RM4001140,
	NT2RM4001200,	NT2RM4001483,	NT2RM4001566,	NT2RM4001592,		
15	NT2RM4001597,	NT2RM4001783,	NT2RM4001823,	NT2RM4001828,	NT2RM4001858,	NT2RM4001979,
	NT2RP1000035,	NT2RP1000111,	NT2RP1000493,	NT2RP1000574,	NT2RP1000630,	NT2RP1000902,
	NT2RP1000915,	NT2RP1000958,	NT2RP1000966,	NT2RP1001013,	NT2RP1001177,	NT2RP2000008,
	NT2RP2000076,	NT2RP2000126,	NT2RP2000153,	NT2RP2000161,	NT2RP2000248,	NT2RP2000258,
	NT2RP2000297,	NT2RP2000420,	NT2RP2000931,	NT2RP2001233,	NT2RP2001420,	NT2RP2001756,
20	NT2RP2001869,	NT2RP2002079,	NT2RP2002270,	NT2RP2002503,	NT2RP2002591,	NT2RP2002880,
	NT2RP2002939,	NT2RP2002993,	NT2RP2003137,	NT2RP2003157,	NT2RP2003277,	NT2RP2003286,
	NT2RP2003308,	NT2RP2003347,	NT2RP2003714,	NT2RP2003912,	NT2RP2004013,	NT2RP2004187,
	NT2RP2004689,	NT2RP2004920,	NT2RP2005393,	NT2RP2005436,	NT2RP2005496,	NT2RP2005539,
	NT2RP2005701,	NT2RP2005767,	NT2RP2005776,	NT2RP2005933,	NT2RP2005942,	NT2RP2006043,
25	NT2RP2006436,	NT2RP3000031,	NT2RP3000050,	NT2RP3000397,	NT2RP3000512,	NT2RP3000527,
	NT2RP3000590,	NT2RP3000603,	NT2RP3000632,	NT2RP3000917,	NT2RP3001057,	NT2RP3001107,
	NT2RP3001120,	NT2RP3001253,	NT2RP3001338,	NT2RP3001384,	NT2RP3001398,	NT2RP3001427,
	NT2RP3001428,	NT2RP3001472,	NT2RP3001646,	NT2RP3001671,	NT2RP3001792,	NT2RP3001855,
	NT2RP3002056,	NT2RP3002165,	NT2RP3002399,	NT2RP3002876,	NT2RP3003193,	NT2RP3003212,
30	NT2RP3003555,	NT2RP3004016,	NT2RP3004206,	NT2RP3004424,	NT2RP3004428,	NT2RP3004566,
	NT2RP3004617,	NT2RP4000078,	NT2RP4000111,	NT2RP4000210,	NT2RP4000398,	NT2RP4000481,
	NT2RP4000518,	NT2RP4000997,	NT2RP4001148,	NT2RP4001206,	NT2RP4001213,	NT2RP4001433,
	NT2RP4001568,	NT2RP4001638,	NT2RP4001696,	NT2RP4001753,	NT2RP4001938,	NT2RP4002058,
	NT2RP4002078,	NT2RP4002081,	NT2RP4002791,	OVARC1000006,	OVARC1000087,	OVARC1000091,
35	OVARC1000241,	OVARC1000326,	OVARC1000556,	OVARC1000846,	OVARC1001038,	OVARC1001180,
	OVARC1001232,	OVARC1001271,	OVARC1001306,	OVARC1001436,	OVARC1002112,	PLACE1000133,
	PLACE1000184,	PLACE1000406,	PLACE1000583,	PLACE1000596,	PLACE1000979,	PLACE1001118,
	PLACE1001383,	PLACE1001632,	PLACE1002171,	PLACE1002433,	PLACE1002438,	PLACE1002532,
	PLACE1002775,	PLACE1002816,	PLACE1002834,	PLACE1003100,	PLACE1003190,	PLACE1003302,
40	PLACE1003519,	PLACE1003521,	PLACE1003605,	PLACE1003704,	PLACE1003738,	PLACE1003885,
	PLACE1003923,	PLACE1004302,	PLACE1004471,	PLACE1004564,	PLACE1004814,	PLACE1004902,
	PLACE1005287,	PLACE1005876,	PLACE1005966,	PLACE1006167,	PLACE1006438,	PLACE1006482,
	PLACE1006829,	PLACE1006878,	PLACE1006917,	PLACE1007014,	PLACE1007547,	PLACE1007598,
	PLACE1007688,	PLACE1007969,	PLACE1008044,	PLACE1008132,	PLACE1008603,	PLACE1009099,
45	PLACE1009130,	PLACE1009308,	PLACE1009398,	PLACE1010134,	PLACE1010194,	PLACE1010702,
	PLACE1010720,	PLACE1010870,	PLACE1011056,	PLACE1011433,	PLACE1011664,	PLACE2000014,
	PLACE2000427,	PLACE3000009,	PLACE3000169,	PLACE4000014,	PLACE4000156,	PLACE4000192,
	PLACE4000261,	PLACE4000326,	PLACE4000489,	SKNMC1000011,	THYRO1000085,	THYRO1000242,
	THYRO1000585,	THYRO1001100,	THYRO1001189,	THYRO1001809,	Y79AA1000037,	Y79AA1000214,
50	Y79AA1000231,	Y79AA1000589,	Y79AA1000752,	Y79AA1001391,	Y79AA1001613,	Y79AA1001705,
	Y79AA1001963,	Y79AA1002431,	Y79AA1002472,	Y79AA1002482		

[0284] The following 292 clones presumably belong to DNA- and/or RNA-binding proteins.

	HEMBA1000158,	HEMBA1000216,	HEMBA1000561,	HEMBA1000591,	HEMBA1000851,	HEMBA1001088,
	HEMBA1001137,	HEMBA1001405,	HEMBA1001510,	HEMBA1001804,	HEMBA1001809,	HEMBA1001819,
55	HEMBA1001847,	HEMBA1001869,	HEMBA1002177,	HEMBA1002935,	HEMBA1003408,	HEMBA1003545,
	HEMBA1003568,	HEMBA1003591,	HEMBA1003662,	HEMBA1003684,	HEMBA1003760,	HEMBA1003783,
	HEMBA1003805,	HEMBA1003953,	HEMBA1004321,	HEMBA1004354,	HEMBA1004389,	HEMBA1004479,
	HEMBA1004669,	HEMBA1004847,	HEMBA1004973,	HEMBA1005202,	HEMBA1005359,	HEMBA1005931,

- PLACE1004804, PLACE1004918, PLACE1005243, PLACE1005305, PLACE1005373, PLACE1005656,
 PLACE1005763, PLACE1005804, PLACE1005953, PLACE1005955, PLACE1006011, PLACE1006469,
 PLACE1006534, PLACE1006626, PLACE1006731, PLACE1006819, PLACE1006829, PLACE1006878,
 PLACE1007226, PLACE1007416, PLACE1007649, PLACE1007706, PLACE1007729, PLACE1007954,
 5 PLACE1007958, PLACE1008111, PLACE1008275, PLACE1008330, PLACE1008643, PLACE1009094,
 PLACE1009130, PLACE1009444, PLACE1009763, PLACE1009861, PLACE1009992, PLACE1009997,
 PLACE1010096, PLACE1010362, PLACE1010481, PLACE1010662, PLACE1011046, PLACE1011219,
 PLACE1011229, PLACE1011332, PLACE1011635, PLACE1011923, PLACE2000021, PLACE2000034,
 PLACE2000398, PLACE2000404, PLACE2000438, PLACE3000009, PLACE3000020, PLACE3000059,
 10 PLACE3000147, PLACE3000339, PLACE3000350, PLACE4000063, PLACE4000100, PLACE4000401,
 PLACE4000548, PLACE4000654, SKNMC1000050, THYRO1000072, THYRO1000197, THYRO1000288,
 THYRO1000605, THYRO1000662, THYRO1000756, THYRO1000852, THYRO1000926, THYRO 1000934,
 THYRO1000951, THYRO 1000983, THYRO1001003, THYRO1001287, THYRO1001374, THYRO1001406,
 THYRO1001617, THYRO1001671, THYRO1001738, Y79AA1000782, Y79AA1001048, Y79AA1001233,
 15 Y79AA1001394, Y79AA1001493, Y79AA1001548, Y79AA1001581, Y79AA1001603, Y79AA1001827,
 Y79AA1002027, Y79AA1002209, Y79AA1002211, Y79AA1002361, Y79AA1002416,
[0282] The following 217 clones presumably belong to a group of cDNAs encoding ATP- and/or GTP-binding proteins.
 HEMBA1000012, HEMBA1000129, HEMBA1000185, HEMBA1000491, HEMBA1000531, HEMBA1001019,
 HEMBA1001174, HEMBA1001387, HEMBA1001595, HEMBA1001723, HEMBA1001913, HEMBA1002161,
 20 HEMBA1002212, HEMBA1002876, HEMBA1002997, HEMBA1003250, HEMBA1003291, HEMBA1003369,
 HEMBA1003555, HEMBA1003560, HEMBA1004131, HEMBA1004199, HEMBA1004202, HEMBA1004354,
 HEMBA1004697, HEMBA1005047, HEMBA1005595, HEMBA1007018, HEMBA1007151, HEMBB1000083,
 HEMBB1000226, HEMBB1000264, HEMBB1000632, HEMBB1000725, HEMBB1001294, HEMBB1002193,
 MAMMA1000085, MAMMA1000612, MAMMA1000731, MAMMA1000738, MAMMA1001038, MAMMA1001735,
 25 MAMMA1001768, MAMMA1003127, NT2RM1000187, NT2RM1000388, NT2RM1000702, NT2RM1000772,
 NT2RM1000924, NT2RM2000469, NT2RM2000577, NT2RM2000740, NT2RM2001100, NT2RM2001201,
 NT2RM2001345, NT2RM2001823, NT2RM2002128, NT2RM4000155, NT2RM4000191, NT2RM4000356,
 NT2RM4000496, NT2RM4000611, NT2RM4000733, NT2RM4000820, NT2RM4001084, NT2RM4001178,
 NT2RM4001344, NT2RM4001444, NT2RM4001592, NT2RM4001714, NT2RM4001758, NT2RM4001880,
 30 NT2RM4002062, NT2RM4002174, NT2RM4002205, NT2RM4002527, NT2RM4002594, NT2RM4002623,
 NT2RP1000470, NT2RP1000478, NT2RP1000915, NT2RP1000958, NT2RP1001080, NT2RP1001410,
 NT2RP1001569, NT2RP2000126, NT2RP2000258, NT2RP2000329, NT2RP2000660, NT2RP2000668,
 NT2RP2000710, NT2RP2000812, NT2RP2000880, NT2RP2001245, NT2RP2001392, NT2RP2002606,
 NT2RP2003277, NT2RP2003912, NT2RP2004538, NT2RP2004568,
 35 NT2RP2004689, NT2RP2004768, NT2RP2004791, NT2RP2004920, NT2RP2005344, NT2RP2005393,
 NT2RP2005763, NT2RP2006534, NT2RP3000046, NT2RP3000252, NT2RP3000350, NT2RP3000359,
 NT2RP3000366, NT2RP3000397, NT2RP3000759, NT2RP3000845, NT2RP3000875, NT2RP3001150,
 NT2RP3001427, NT2RP3001453, NT2RP3001529, NT2RP3001730, NT2RP3001799, NT2RP3001857,
 NT2RP3001938, NT2RP3002007, NT2RP3002151, NT2RP3002330, NT2RP3002399, NT2RP3002671,
 40 NT2RP3003301, NT2RP3003353, NT2RP3003589, NT2RP3003809, NT2RP3003876, NT2RP3004189,
 NT2RP3004428, NT2RP3004578, NT2RP4000290, NT2RP4000481, NT2RP4000518, NT2RP4000781,
 NT2RP4000839, NT2RP4000929, NT2RP4001041, NT2RP4001079, NT2RP4001375, NT2RP4001414,
 NT2RP4001592, NT2RP4001634, NT2RP4001644, NT2RP4001656, NT2RP4001896, NT2RP4002047,
 NT2RP4002058, NT2RP4002408, NT2RP5003477, OVARC1000013, OVARC1000304, OVARC1000556,
 45 OVARC1000771, OVARC1000800, OVARC1001068, OVARC1002138, PLACE1000040, PLACE1000588,
 PLACE1001104, PLACE1001739, PLACE1002433, PLACE1002437, PLACE1002714, PLACE1003394,
 PLACE1003521, PLACE1003915, PLACE1004902, PLACE1005243, PLACE1005305, PLACE1005549,
 PLACE1005739, PLACE1005921, PLACE1006119, PLACE1006196, PLACE1006552, PLACE1006956,
 PLACE1007409, PLACE1007697, PLACE1007946, PLACE1008244, PLACE1009404, PLACE1009476,
 50 PLACE1009596, PLACE1009908, PLACE1010134, PLACE1010720, PLACE1010896, PLACE1011109,
 PLACE1011114, PLACE1011310, PLACE1011922, PLACE2000014, PLACE2000039, PLACE2000274,
 PLACE2000404, PLACE2000427, PLACE3000350, PLACE4000009, PLACE4000014, PLACE4000326,
 SKNMC1000013, THYRO1000072, THYRO1001458, Y79AA1000833, Y79AA1000962, Y79AA1001394,
 Y79AA1001875, Y79AA1001963, Y79AA1002209,
 55 **[0283]** The following 320 clones presumably belong to nuclear proteins.
 HEMBA1000005, HEMBA1000158, HEMBA1000216, HEMBA1000561, HEMBA1000591, HEMBA1001088,
 HEMBA1001137, HEMBA1001405, HEMBA1001510, HEMBA1001579, HEMBA1001809, HEMBA1001819,
 HEMBA1001824, HEMBA1001847, HEMBA1001869, HEMBA1002177, HEMBA1002241, HEMBA1002495,

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Y79AA1001874(600315), Y79AA1002204(605033), Y79AA1002210(191161), Y79AA1002472(603971),
Y79AA1002482(603971),

[0281] The following 425 clones presumably belong to enzymes and/or metabolism-associated proteins.

	HEMBA1000012,	HEMBA1000129,	HEMBA1000141,	HEMBA1000150,	HEMBA1000542,	HEMBA1000852,
5	HEMBA1001019,	HEMBA1001257,	HEMBA1001526,	HEMBA1001620,	HEMBA1001866,	HEMBA1001896,
	HEMBA1002212,	HEMBA1002513,	HEMBA1002746,	HEMBA1002973,	HEMBA1003046,	HEMBA1003136,
	HEMBA1003179,	HEMBA1003250,	HEMBA1003291,	HEMBA1003408,	HEMBA1003538,	HEMBA1003679,
	HEMBA1003680,	HEMBA1004199,	HEMBA1004227,	HEMBA1004408,	HEMBA1004509,	HEMBA1004734,
	HEMBA1004768,	HEMBA1005394,	HEMBA1005513,	HEMBA1005737,	HEMBA1005815,	HEMBA1006031,
10	HEMBA1006272,	HEMBA1006278,	HEMBA1006291,	HEMBA1006309,	HEMBA1006347,	HEMBA1006485,
	HEMBA1006521,	HEMBA1006624,	HEMBA1006885,	HEMBA1006976,	HEMBA1007121,	HEMBA1007224,
	HEMBA1007243,	HEMBA1007300,	HEMBA1000083,	HEMBA1000217,	HEMBA1000915,	HEMBA1000947,
	HEMBA1001137,	HEMBA1001346,	HEMBA1001429,	HEMBA1001443,	HEMBA1001915,	HEMBA1001950,
	HEMBA1002042,	MAMMA1000020,	MAMMA1000085,	MAMMA1000672,	MAMMA1000713,	MAMMA1000841,
15	MAMMA1000897,	MAMMA1001008,	MAMMA1001038,	MAMMA1001059,	MAMMA1001476,	MAMMA1001501,
	MAMMA1002268,	MAMMA1002470,	MAMMA1002530,	MAMMA1002573,	MAMMA1002619,	MAMMA1002655,
	MAMMA1002671,	MAMMA1003013,	MAMMA1003035,	NT2RM1000039,	NT2RM1000132,	NT2RM1000153,
	NT2RM1000256,	NT2RM1000280,	NT2RM1000377,	NT2RM1000553,	NT2RM1000648,	NT2RM1000702,
	NT2RM1000894,	NT2RM1001072,	NT2RM1001115,	NT2RM2000013,	NT2RM2000092,	NT2RM2000322,
20	NT2RM2000368,	NT2RM2000371,	NT2RM2000469,	NT2RM2000504,	NT2RM2000577,	NT2RM2000594,
	NT2RM2000951,	NT2RM2001238,	NT2RM2001547,	NT2RM2001632,	NT2RM2001664,	NT2RM2001698,
	NT2RM2001700,	NT2RM2001730,	NT2RM2001782,	NT2RM2001803,	NT2RM2001886,	NT2RM2001935,
	NT2RM2001997,	NT2RM2002030,	NT2RM2002128,	NT2RM4000024,	NT2RM4000155,	NT2RM4000344,
	NT2RM4000471,	NT2RM4000616,	NT2RM4000657,	NT2RM4000712,	NT2RM4000820,	NT2RM4001313,
25	NT2RM4001316,	NT2RM4001444,	NT2RM4001592,	NT2RM4001758,	NT2RM4001819,	NT2RM4001880,
	NT2RM4002062,	NT2RM4002063,	NT2RM4002189,	NT2RM4002213,	NT2RM4002251,	NT2RM4002409,
	NT2RM4002532,	NT2RM4002623,	NT2RP1000376,	NT2RP1000443,	NT2RP1000522,	NT2RP1000834,
	NT2RP1000947,	NT2RP1001079,	NT2RP1001185,	NT2RP1001253,	NT2RP1001361,	NT2RP1001543,
	NT2RP2000056,	NT2RP2000114,	NT2RP2000183,	NT2RP2000248,	NT2RP2000329,	NT2RP2000422,
30	NT2RP2000448,	NT2RP2000668,	NT2RP2000710,	NT2RP2000816,	NT2RP2001070,	NT2RP2001392,
	NT2RP2001601,	NT2RP2001663,	NT2RP2001740,	NT2RP2001748,	NT2RP2001898,	NT2RP2002124,
	NT2RP2002256,	NT2RP2002609,	NT2RP2002618,	NT2RP2002959,	NT2RP2002993,	NT2RP2003230,
	NT2RP2003286,	NT2RP2003401,	NT2RP2003506,	NT2RP2003543,	NT2RP2003643,	NT2RP2003702,
	NT2RP2003704,	NT2RP2003713,	NT2RP2003737,	NT2RP2003840,	NT2RP2003912,	NT2RP2003952,
35	NT2RP2004098,	NT2RP2004239,	NT2RP2004245,	NT2RP2004768,	NT2RP2004791,	NT2RP2004799,
	NT2RP2004933,	NT2RP2005038,	NT2RP2005139,	NT2RP2005162,	NT2RP2005204,	NT2RP2005239,
	NT2RP2005276,	NT2RP2005344,	NT2RP2005360,	NT2RP2005457,	NT2RP2005498,	NT2RP2005549,
	NT2RP2005557,	NT2RP2005605,	NT2RP2005635,	NT2RP2005723,	NT2RP2005773,	NT2RP2005775,
	NT2RP2005776,	NT2RP2005784,	NT2RP2005835,	NT2RP2005942,	NT2RP2006534,	NT2RP2006571,
40	NT2RP2006573,	NT2RP3000031,	NT2RP3000085,	NT2RP3000207,	NT2RP3000359,	NT2RP3000578,
	NT2RP3000742,	NT2RP3000845,	NT2RP3000875,	NT2RP3000917,	NT2RP3001055,	NT2RP3001221,
	NT2RP3001495,	NT2RP3001898,	NT2RP3001938,	NT2RP3002303,	NT2RP3002351,	NT2RP3002501,
	NT2RP3002602,	NT2RP3002628,	NT2RP3002663,	NT2RP3003301,	NT2RP3003385,	NT2RP3003490,
	NT2RP3003659,	NT2RP3003825,	NT2RP3003831,	NT2RP3003846,	NT2RP3003914,	NT2RP3004148,
45	NT2RP3004209,	NT2RP3004378,	NT2RP3004669,	NT2RP3004670,	NT2RP4000259,	NT2RP4000312,
	NT2RP4000367,	NT2RP4000417,	NT2RP4000457,	NT2RP4000657,	NT2RP4000817,	NT2RP4000855,
	NT2RP4000879,	NT2RP4000927,	NT2RP4000973,	NT2RP4000997,	NT2RP4001041,	NT2RP4001079,
	NT2RP4001095,	NT2RP4001143,	NT2RP4001219,	NT2RP4001375,	NT2RP4001389,	NT2RP4001483,
	NT2RP4001555,	NT2RP4001592,	NT2RP4001644,	NT2RP4001730,	NT2RP4001946,	NT2RP4002408,
50	NT2RP5003500,	NT2RP5003522,	OVARC1000013,	OVARC1000060,	OVARC1000139,	OVARC1000288,
	OVARC1000309,	OVARC1000473,	OVARC1000556,	OVARC1000682,	OVARC1000722,	OVARC1000751,
	OVARC1000885,	OVARC1000915,	OVARC1001107,	OVARC1001713,	OVARC1001762,	OVARC1001809,
	OVARC1001942,	OVARC1002156,	OVARC1002165,	PLACE1000007,	PLACE1000142,	PLACE1000185,
	PLACE1000213,	PLACE1000383,	PLACE1000420,	PLACE1000547,	PLACE1000653,	PLACE1000755,
55	PLACE1001054,	PLACE1001062,	PLACE1001672,	PLACE1001692,	PLACE1001748,	PLACE1001781,
	PLACE1001817,	PLACE1001869,	PLACE1001989,	PLACE1002073,	PLACE1002598,	PLACE1002908,
	PLACE1002991,	PLACE1003174,	PLACE1003176,	PLACE1003709,	PLACE1003885,	PLACE1003888,
	PLACE1003903,	PLACE1003915,	PLACE1004270,	PLACE1004428,	PLACE1004437,	PLACE1004751,

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	NT2RP2000248(603367), NT2RP2000420(600834), NT2RP2001233(603971), NT2RP2001397(602755),	NT2RP2000297(602277), NT2RP2000523(600130), NT2RP2001327(191161), NT2RP2001460(190370),	NT2RP2000310(239500), NT2RP2000809(603885), NT2RP2001378(158370), NT2RP2001520(603667),	NT2RP2000414(601037), NT2RP2000812(160777), NT2RP2001394(300208), NT2RP2001536(600675),
5	NT2RP2001876(601833), NT2RP2002124(603486), NT2RP2003000(191161), NT2RP2003295(603494), NT2RP2003714(603971),	NT2RP2001898(147264), NT2RP2002325(603866), NT2RP2003157(601940), NT2RP2003517(190040), NT2RP2003737(602962),	NT2RP2002025(601581), NT2RP2002503(601781), NT2RP2003164(604746), NT2RP2003564(109092), NT2RP2003952(602675),	NT2RP2002058(604737), NT2RP2002959(602962), NT2RP2003228(602638), NT2RP2003604(604785), NT2RP2004013(602542),
10	NT2RP2004170(300196), NT2RP2005003(109092), NT2RP2005288(603524), NT2RP2005358(603573), NT2RP2005525(602655),	NT2RP2004587(162250), NT2RP2005144(604730), NT2RP2005315(604039), NT2RP2005407(167040), NT2RP2005719(601178),	NT2RP2004732(162250), NT2RP2005239(603485), NT2RP2005325(603759), NT2RP2005436(601940), NT2RP2006043(601940),	NT2RP2004933(603289), NT2RP2005276(602371), NT2RP2005336(190370), NT2RP2005476(602680), NT2RP2006071(604299),
15	NT2RP2006219(601279), NT2RP3000068(182530), NT2RP3000596(190370), NT2RP3001081(603524), NT2RP3001428(189940),	NT2RP2006312(603111), NT2RP3000085(300032), NT2RP3000739(125370), NT2RP3001216(603121), NT2RP3001723(604569),	NT2RP2006456(604619), NT2RP3000299(602941), NT2RP3000753(162230), NT2RP3001307(180069), NT2RP3001855(602100),	NT2RP3000050(603971), NT2RP3000403(604981), NT2RP3001057(603971), NT2RP3001338(314998), NT2RP3001898(604561),
20	NT2RP3001969(190370), NT2RP3002351(604887), NT2RP3003251(109092), NT2RP3003672(313470), NT2RP3004209(125255),	NT2RP3002056(180201), NT2RP3002399(602638), NT2RP3003282(602378), NT2RP3003831(604051), NT2RP3004258(604347),	NT2RP3002062(603885), NT2RP3002953(604967), NT2RP3003313(603810), NT2RP3004016(601742), NT2RP3004490(603328),	NT2RP3002151(139259), NT2RP3002988(603258), NT2RP3003409(604533), NT2RP3004078(142765), NT2RP3004534(600586),
25	NT2RP3004569(106410), NT2RP4000376(603873), NT2RP4000925(600245), NT2RP4001213(602277), NT2RP4001483(203740),	NT2RP3004572(604912), NT2RP4000449(604479), NT2RP4001086(162230), NT2RP4001276(190370), NT2RP4001575(603443),	NT2RP4000109(603745), NT2RP4000855(602675), NT2RP4001126(190370), NT2RP4001407(190370), NT2RP4001760(305400),	NT2RP4000367(603722), NT2RP4000879(314370), NT2RP4001150(601581), NT2RP4001433(602277), NT2RP4001861(190370),
30	NT2RP4002078(603971), OVARC1000520(604126), OVARC1001113(602121), OVARC1001496(602619), OVARC1001766(603910),	NT2RP4002791(189940), OVARC1000722(604014), OVARC1001244(601540), OVARC1001506(601313), OVARC1001809(603730),	OVARC1000014(603371), OVARC1000771(179509), OVARC1001372(603145), OVARC1001577(603269), PLACE1000133(602542),	OVARC1000139(603486), OVARC1001051(600051), OVARC1001417(300182), OVARC1001726(300103), PLACE1000383(300171),
35	PLACE1000420(600312), PLACE1001387(600206), PLACE1002437(600046), PLACE1002834(194558), PLACE1003353(604704),	PLACE1000583(194558), PLACE1001602(604913), PLACE1002474(602108), PLACE1002908(604327), PLACE1003366(603681),	PLACE1000588(600411), PLACE1002046(151625), PLACE1002685(604515), PLACE1003045(173910), PLACE1003493(601456),	PLACE1001171(310400), PLACE1002140(603748), PLACE1002782(602095), PLACE1003302(194558), PLACE1003669(190370),
40	PLACE1003704(601940), PLACE1004197(601610), PLACE1004471(194558), PLACE1004777(118423), PLACE1006170(601026),	PLACE1003903(123860), PLACE1004277(603493), PLACE1004506(603450), PLACE1004814(601940), PLACE1006438(600834),	PLACE1003968(602742), PLACE1004316(604261), PLACE1004510(604912), PLACE1005494(603652), PLACE1006615(603910),	PLACE1004183(604701), PLACE1004358(603272), PLACE1004674(601057), PLACE1006040(603061), PLACE1007140(190370),
45	PLACE1007239(604784), PLACE1008177(190370), PLACE1009027(300121), PLACE1009524(602488), PLACE1011635(604058),	PLACE1007257(300108), PLACE1008356(604039), PLACE1009113(600675), PLACE1010529(604834), PLACE1011858(603882),	PLACE1007511(148020), PLACE1008402(603344), PLACE1009158(604140), PLACE1010870(603971), PLACE1011922(160776),	PLACE1007598(602277), PLACE1008696(602141), PLACE1009444(600286), PLACE1010896(160776), PLACE2000015(600051),
50	PLACE2000072(603430), PLACE2000458(600976), PLACE4000156(603971), THYRO1000034(190370), THYRO1000501(109092),	PLACE2000216(182790), PLACE3000242(300132), PLACE4000369(603808), THYRO1000327(603243), THYRO1000662(278750),	PLACE2000399(313470), PLACE4000009(160776), SKNMC1000046(603144), THYRO1000343(125370), THYRO1000684(603885),	PLACE2000438(602273), PLACE4000014(300032), SKNMC1000050(114230), THYRO1000358(604188), THYRO1000748(300023),
55	THYRO1000934(179035), THYRO1001458(160776), Y79AA1000469(602434), Y79AA1001391(142959),	THYRO1001120(602582), THYRO1001617(602744), Y79AA1000560(601026), Y79AA1001548(600286),	THYRO1001189(603971), THYRO1001671(603281), Y79AA1000734(603867), Y79AA1001594(600936),	THYRO1001204(603169), Y79AA1000346(604355), Y79AA1000782(600417), Y79AA1001711(600063),

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	NT2RP2005436,	NT2RP2005476,	NT2RP2005525,	NT2RP2005694,	NT2RP2005719,	NT2RP2006043,
	NT2RP2006071,	NT2RP2006219,	NT2RP2006312,	NT2RP2006456,	NT2RP3000050,	NT2RP3000068,
	NT2RP3000085,	NT2RP3000299,	NT2RP3000403,	NT2RP3000596,	NT2RP3000739,	NT2RP3000753,
5	NT2RP3000875,	NT2RP3001057,	NT2RP3001081,	NT2RP3001216,	NT2RP3001307,	NT2RP3001338,
	NT2RP3001427,	NT2RP3001428,	NT2RP3001679,	NT2RP3001723,	NT2RP3001855,	NT2RP3001898,
	NT2RP3001969,	NT2RP3002056,	NT2RP3002062,	NT2RP3002151,	NT2RP3002351,	NT2RP3002399,
	NT2RP3002953,	NT2RP3002988,	NT2RP3003078,	NT2RP3003251,	NT2RP3003282,	NT2RP3003313,
	NT2RP3003327,	NT2RP3003409,	NT2RP3003672,	NT2RP3003831,	NT2RP3004016,	NT2RP3004078,
10	NT2RP3004209,	NT2RP3004258,	NT2RP3004490,	NT2RP3004534,	NT2RP3004569,	NT2RP3004572,
	NT2RP4000109,	NT2RP4000367,	NT2RP4000376,	NT2RP4000449,	NT2RP4000855,	NT2RP4000879,
	NT2RP4000925,	NT2RP4001086,	NT2RP4001126,	NT2RP4001150,	NT2RP4001213,	NT2RP4001276,
	NT2RP4001407,	NT2RP4001433,	NT2RP4001483,	NT2RP4001575,	NT2RP4001760,	NT2RP4001861,
	NT2RP4002078,	NT2RP4002791,	OVARC1000014,	OVARC1000139,	OVARC1000520,	OVARC1000722,
15	OVARC1000771,	OVARC1000834,	OVARC1001051,	OVARC1001113,	OVARC1001244,	OVARC1001372,
	OVARC1001417,	OVARC1001496,	OVARC1001506,	OVARC1001577,	OVARC1001726,	OVARC1001766,
	OVARC1001809,	OVARC1002165,	PLACE1000133,	PLACE1000383,	PLACE1000420,	PLACE1000583,
	PLACE1000588,	PLACE1001171,	PLACE1001387,	PLACE1001602,	PLACE1002046,	PLACE1002140,
	PLACE1002437,	PLACE1002474,	PLACE1002685,	PLACE1002782,	PLACE1002834,	PLACE1002908,
20	PLACE1003045,	PLACE1003302,	PLACE1003353,	PLACE1003366,	PLACE1003493,	PLACE1003669,
	PLACE1003704,	PLACE1003903,	PLACE1003968,	PLACE1004183,	PLACE1004197,	PLACE1004277,
	PLACE1004316,	PLACE1004358,	PLACE1004471,	PLACE1004506,	PLACE1004510,	PLACE1004674,
	PLACE1004777,	PLACE1004814,	PLACE1005494,	PLACE1006040,	PLACE1006170,	PLACE1006438,
	PLACE1006615,	PLACE1007140,	PLACE1007239,	PLACE1007257,	PLACE1007511,	PLACE1007598,
25	PLACE1008177,	PLACE1008356,	PLACE1008402,	PLACE1008696,	PLACE1009027,	PLACE1009113,
	PLACE1009158,	PLACE1009444,	PLACE1009524,	PLACE1010529,	PLACE1010870,	PLACE1010896,
	PLACE1011635,	PLACE1011858,	PLACE1011922,	PLACE2000015,	PLACE2000072,	PLACE2000216,
	PLACE2000399,	PLACE2000438,	PLACE2000458,	PLACE3000242,	PLACE4000009,	PLACE4000014,
	PLACE4000156,	PLACE4000369,	SKNMC1000046,	SKNMC1000050,	THYRO1000034,	THYRO1000327,
30	THYRO1000343,	THYRO1000358,	THYRO1000501,	THYRO1000662,	THYRO1000684,	THYRO1000748,
	THYRO1000934,	THYRO1001120,	THYRO1001189,	THYRO1001204,	THYRO1001458,	THYRO1001617,
	THYRO1001671,	Y79AA1000346,	Y79AA1000469,	Y79AA1000560,	Y79AA1000734,	Y79AA1000782,
	Y79AA1001391,	Y79AA1001548,	Y79AA1001594,	Y79AA1001711,	Y79AA1001874,	Y79AA1002204,
	Y79AA1002210,	Y79AA1002258,	Y79AA1002472,	Y79AA1002482,		
35	[0280] Among them, Swiss-Prot database search and GenBank or UniGene database search revealed that the following 380 clones matched the data of genes or proteins which had been registered in the database of Online Mendelian Inheritance in Man (OMIM) (http://www.ncbi.nlm.nih.gov/omim/), which is a database of human genes and diseases. (The corresponding OMIM numbers are parenthetically indicated following the clone names.)					
	HEMBB1000985(147485),	HEMBB1001068(603142),	HEMBB1001282(182900),	HEMBB1001339(300080),		
40	HEMBB1001482(603971),	HEMBB1001564(603931),	HEMBB1001802(125660),	HEMBB1001905(190370),		
	HEMBB1001908(601408),	HEMBB1002217(603971),	HEMBB1002477(604439),	MAMMA1000388(604865),		
	MAMMA1000731(602118),	MAMMA1001305(602732),	MAMMA1001633(600834),	MAMMA1001868(190370),		
	MAMMA1002170(603624),	MAMMA1002198(600538),	MAMMA1002268(603730),	MAMMA1002485(603665),		
	MAMMA1002530(603602),	MAMMA1002858(601064),	MAMMA1002869(602567),	MAMMA1002881(602692),		
45	MAMMA1003047(603566),	MAMMA1003146(603094),	MAMMA1003166(604061),	NT2RM1000001(601169),		
	NT2RM1000153(600417),	NT2RM1000252(604108),	NT2RM1000555(191510),	NT2RM1000770(300061),		
	NT2RM1000826(191510),	NT2RM1000850(182900),	NT2RM1001003(604785),	NT2RM1001092(603971),		
	NT2RM1001102(604533),	NT2RM2000191(602973),	NT2RM2000363(151410),	NT2RM2000594(602900),		
	NT2RM2000624(601940),	NT2RM2000714(179555),	NT2RM2000821(600959),	NT2RM2001035(604913),		
50	NT2RM2001575(109092),	NT2RM2001652(604141),	NT2RM2001664(603722),	NT2RM2001668(602952),		
	NT2RM2001698(604327),	NT2RM2001803(603722),	NT2RM2001839(603420),	NT2RM4000155(187790),		
	NT2RM4000471(603485),	NT2RM4000486(168730),	NT2RM4000657(602142),	NT2RM4000751(602277),		
	NT2RM4000996(603971),	NT2RM4001629(601114),	NT2RM4001810(155760),	NT2RM4001819(176873),		
	NT2RM4001876(179555),	NT2RM4001940(603887),	NT2RM4002066(300188),	NT2RM4002093(600693),		
55	NT2RM4002146(602603),	NT2RM4002161(254780),	NT2RM4002558(604194),	NT2RM4002571(602274),		
	NT2RP1000086(602219),	NT2RP1000574(601740),	NT2RP1000825(602732),	NT2RP1000833(602973),		
	NT2RP1000959(180510),	NT2RP1000966(164035),	NT2RP1001013(194558),	NT2RP1001185(243500),		
	NT2RP1001482(600586),	NT2RP1001665(114180),	NT2RP2000070(600976),	NT2RP2000147(603535),		

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	NT2RP3000320, NT2RP3000632, NT2RP3001268, NT2RP3002165, NT2RP3003313, NT2RP3004428, NT2RP4000455, NT2RP4001080, NT2RP4001638, NT2RP4002078, OVARC1001271, PLACE1000786, PLACE1001383, PLACE1002532, PLACE1003885, PLACE1006318, PLACE1007547, PLACE1009246, PLACE1010771, PLACE3000169, PLACE4000261, THYRO1000488, Y79AA1000013, Y79AA1001299, Y79AA1002103,	NT2RP3000512, NT2RP3001057, NT2RP3001338, NT2RP3002399, NT2RP3003327, NT2RP3004498, NT2RP4000648, NT2RP4001213, NT2RP4001753, NT2RP4002081, OVARC1001417, PLACE1000979, PLACE1001602, PLACE1002775, PLACE1004471, PLACE1006438, PLACE1007598, PLACE1009308, PLACE1010870, PLACE3000254, PLACE4000450, THYRO1000501, Y79AA1000033, Y79AA1001312,	NT2RP3000527, NT2RP3001107, NT2RP3001398, NT2RP3002876, NT2RP3003555, NT2RP3004566, NT2RP4000837, NT2RP4001433, NT2RP4001760, NT2RP5003461, OVARC1001436, PLACE1001118, PLACE1001632, PLACE1002834, PLACE1005584, PLACE1006482, PLACE1007955, PLACE1009398, PLACE1011160, PLACE4000128, PLACE4000489, THYRO1000569, Y79AA1000037, Y79AA1001391,	NT2RP3000590, NT2RP3001111, NT2RP3001527, NT2RP3003133, NT2RP3004016, NT2RP3004617, NT2RP4000865, NT2RP4001529, NT2RP4001790, OVARC1000151, PLACE1000133, PLACE1001238, PLACE1002171, PLACE1003302, PLACE1005803, PLACE1007239, PLACE1008132, PLACE1009798, PLACE1011433, PLACE4000156, THYRO1000085, THYRO 1001100, Y79AA1000342, Y79AA1001533,	NT2RP3000603, NT2RP3001120, NT2RP3001688, NT2RP3003193, NT2RP3004125, NT2RP4000210, NT2RP4000997, NT2RP4001551, NT2RP4001838, OVARC1000241, PLACE1000583, PLACE1001294, PLACE1002438, PLACE1003605, PLACE1005966, PLACE1007346, PLACE1008201, PLACE1010134, PLACE1011576, PLACE4000192, THYRO1000121, THYRO1001189, Y79AA1000627, Y79AA1001613,	NT2RP3000605, NT2RP3001150, NT2RP3001855, NT2RP3003251, NT2RP3004242, NT2RP4000398, NT2RP4001029, NT2RP4001568, NT2RP4001938, OVARC1000479, PLACE1000706, PLACE1001304, PLACE1002450, PLACE1003738, PLACE1006167, PLACE1007488, PLACE1009099, PLACE1010702, PLACE3000009, PLACE4000211, THYRO1000242, THYRO1001809, Y79AA1000705, Y79AA1001866,
	[0279] The following 392 clones are categorized into disease-associated proteins.					
	HEMBA 1000020, HEMBA 1001043, HEMBA 1001672, HEMBA 1002547, HEMBA 1003369, HEMBA 1003568, HEMBA 1004275, HEMBA 1004669, HEMBA 1005581, HEMBA 1006737, HEMBA 100693, HEMBA 1001482, HEMBA 1002477, MAMMA 1002170, MAMMA 1002869, NT2RM 1000153, NT2RM 1001003,	HEMBA 1000216, HEMBA 1001059, HEMBA 1001819, HEMBA 1002555, HEMBA 1003417, HEMBA 1003569, HEMBA 1004321, HEMBA 1005009, HEMBA 1005621, HEMBA 1006807, HEMBA 1000927, HEMBA 1001564, MAMMA 1000388, MAMMA 1002198, MAMMA 1002881, NT2RM 1000252, NT2RM 1001092,	HEMBA 1000304, HEMBA 1001071, HEMBA 1001921, HEMBA 1002810, HEMBA 1003418, HEMBA 1003581, HEMBA 1004353, HEMBA 1005338, HEMBA 1005699, HEMBA 1006877, HEMBA 1000985, HEMBA 1001802, MAMMA 1000731, MAMMA 1002268, MAMMA 1003047, NT2RM 1000555, NT2RM 1001102,	HEMBA 1000561, HEMBA 1001088, HEMBA 1002267, HEMBA 1002939, HEMBA 1003433, HEMBA 1004168, HEMBA 1004356, HEMBA 1005367, HEMBA 1006507, HEMBA 1007121, HEMBA 1001068, HEMBA 1001905, MAMMA 1001305, MAMMA 1002485, MAMMA 1003146, NT2RM 1000770, NT2RM 2000191,	HEMBA 1000569, HEMBA 1001569, HEMBA 1002419, HEMBA 1002997, HEMBA 1003538, HEMBA 1004202, HEMBA 1004479, HEMBA 1005423, HEMBA 1006650, HEMBA 1007243, HEMBA 1001282, HEMBA 1001908, MAMMA 1001633, MAMMA 1002530, MAMMA 1003166, NT2RM 1000826, NT2RM 2000821,	HEMBA 1000910, HEMBA 1001661, HEMBA 1002469, HEMBA 1003148, HEMBA 1003555, HEMBA 1004248, HEMBA 1004509, HEMBA 1005528, HEMBA 1006652, HEMBA 1000119, HEMBA 1001339, HEMBA 1002217, MAMMA 1001868, MAMMA 1002858, NT2RM 1000001, NT2RM 1000850,
	NT2RM 1001003, NT2RM 2000363, NT2RM 2001575, NT2RM 2001839, NT2RM 4000996, NT2RM 4001940, NT2RM 4002558, NT2RP 1000833, NT2RP 1001665, NT2RP 2000310, NT2RP 2001233, NT2RP 2001520, NT2RP 2002124, NT2RP 2003164, NT2RP 2003714, NT2RP 2004732,	NT2RM 2000594, NT2RM 2001652, NT2RM 4000155, NT2RM 4001629, NT2RM 4002066, NT2RM 4002571, NT2RP 1000959, NT2RP 2000070, NT2RP 2000414, NT2RP 2001327, NT2RP 2001536, NT2RP 2002325, NT2RP 2003228, NT2RP 2003737, NT2RP 2004933,	NT2RM 2000624, NT2RM 2001664, NT2RM 4000471, NT2RM 4001810, NT2RM 4002093, NT2RP 1000086, NT2RP 1000966, NT2RP 2000147, NT2RP 2000420, NT2RP 2001378, NT2RP 2001876, NT2RP 2002503, NT2RP 2003295, NT2RP 2003952, NT2RP 2005003,	NT2RM 2000714, NT2RM 2001668, NT2RM 4000486, NT2RM 4001819, NT2RM 4002146, NT2RP 1000574, NT2RP 1001013, NT2RP 2000224, NT2RP 2000523, 		

PLACE1002722, PLACE1003045, PLACE1003238, PLACE1003258, PLACE1003493, PLACE1004197,
 PLACE1004793, PLACE1005953, PLACE1005955, PLACE1006157, PLACE1006239, PLACE1006368,
 PLACE1006534, PLACE1006754, PLACE1006956, PLACE1007416, PLACE1007632, PLACE1007649,
 PLACE1008643, PLACE1009094, PLACE1009992, PLACE1010231, PLACE1010662, PLACE1011371,
 5 PLACE2000034, PLACE2000373, PLACE2000398, PLACE2000399, PLACE2000438, PLACE2000458,
 PLACE3000339, PLACE4000063, PLACE4000230, PLACE4000522, PLACE4000548, PLACE4000581,
 THYRO1000327, THYRO1000756, THYRO1001287, Y79AA1001603, Y79AA1001874

[0277] The following 129 clones are categorized into signal transduction-associated proteins.

10 HEMBA1000303, HEMBA1000369, HEMBA1000608, HEMBA1000657, HEMBA1000919, HEMBA1001019,
 HEMBA1001174, HEMBA1001822, HEMBA1001921, HEMBA1002139, HEMBA1002212, HEMBA1002341,
 HEMBA1002417, HEMBA1002768, HEMBA1003250, HEMBA1003291, HEMBA1003645, HEMBA1004286,
 HEMBA1005737, HEMBA1006130, HEMBA1006708, HEMBB1000083, HEMBB1000266, HEMBB1000632,
 HEMBB1000781, HEMBB1000831, HEMBB1002193, MAMMA1000173, MAMMA1001038, MAMMA1001198,
 MAMMA1002842, MAMMA1003057, NT2RM1000702, NT2RM1000772, NT2RM1001072, NT2RM2000030,
 15 NT2RM2000469, NT2RM2000612, NT2RM2001221, NT2RM2001345, NT2RM2002128, NT2RM4000229,
 NT2RM4000354, NT2RM4000611, NT2RM4000798, NT2RM4001411, NT2RM4001629, NT2RM4001294,
 NT2RM4001758, NT2RM4002013, NT2RM4002527, NT2RP1000018, NT2RP1000701, NT2RP1001294,
 NT2RP1001302, NT2RP2000668, NT2RP2001440, NT2RP2001560, NT2RP2002058, NT2RP2002193,
 NT2RP2002408, NT2RP2002710, NT2RP2002929, NT2RP2003164, NT2RP2003912, NT2RP2004232,
 20 NT2RP2004768, NT2RP2006071, NT2RP2006534, NT2RP3000759, NT2RP3000845, NT2RP3001646,
 NT2RP3001857, NT2RP3001938, NT2RP3002004, NT2RP3002785, NT2RP3002909, NT2RP3002988,
 NT2RP3003800, NT2RP3004189, NT2RP3004544, NT2RP4000147, NT2RP4000839, NT2RP4001122,
 NT2RP4001148, NT2RP4001336, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001849,
 NT2RP4001896, NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000013, OVARC1000437,
 25 OVARC1000556, OVARC1000649, OVARC 1000945, OVARC1001200, OVARC1002182, PLACE1000977,
 PLACE1001387, PLACE1002493, PLACE1002591, PLACE1003190, PLACE1003353, PLACE1004128,
 PLACE1004302, PLACE1004937, PLACE1005243, PLACE1008000, PLACE1008244, PLACE1008650,
 PLACE1009468, PLACE1009596, PLACE1009708, PLACE1009845, PLACE1010926, PLACE1011041,
 PLACE2000164, PLACE2000371, PLACE3000145, PLACE3000350, THYRO1000072, THYRO1000748,
 30 THYRO1001120, Y79AA1000328, Y79AA1002431

[0278] The following 309 clones are categorized into transcription-associated proteins.

HEMBA1000158, HEMBA1000201, HEMBA1000216, HEMBA1000555, HEMBA1000561, HEMBA1000851,
 HEMBA1001077, HEMBA1001137, HEMBA1001405, HEMBA1001510, HEMBA1001635, HEMBA1001804,
 HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1001869, HEMBA1002035, HEMBA1002092,
 35 HEMBA1002177, HEMBA1002770, HEMBA1002935, HEMBA1003408, HEMBA1003545, HEMBA1003568,
 HEMBA1003662, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321,
 HEMBA1004353, HEMBA1004389, HEMBA1004479, HEMBA1004758, HEMBA1004973, HEMBA1005219,
 HEMBA1005359, HEMBA1005513, HEMBA1005528, HEMBA1005548, HEMBA1005558, HEMBA1005931,
 HEMBA1006158, HEMBA1006248, HEMBA1006278, HEMBA1006283, HEMBA1006347, HEMBA1006359,
 40 HEMBA1006559, HEMBA1006941, HEMBB1000789, HEMBB1001011, HEMBB1001314, HEMBB1001482,
 HEMBB1001673, HEMBB1001749, HEMBB1001839, HEMBB1001908, HEMBB1002134, HEMBB1002217,
 HEMBB1002342, HEMBB1002607, MAMMA1000183, MAMMA1000388, MAMMA1001105, MAMMA1001222,
 MAMMA1001260, MAMMA1001627, MAMMA1001633, MAMMA1001743, MAMMA1001820, MAMMA1001837,
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 45 NT2RM1000885, NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000452, NT2RM2000735,
 NT2RM2000740, NT2RM2001035, NT2RM2001105, NT2RM2001575, NT2RM2001670, NT2RM2001716,
 NT2RM2001771, NT2RM2002091, NT2RM4000024, NT2RM4000046, NT2RM4000104, NT2RM4000202,
 NT2RM4000531, NT2RM4000595, NT2RM4000733, NT2RM4000734,
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 50 NT2RM4001483, NT2RM4001592, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858,
 NT2RM4001979, NT2RM4002066, NT2RP1000086, NT2RP1000111, NT2RP1000574, NT2RP1000902,
 NT2RP1001013, NT2RP2000008, NT2RP2000126, NT2RP2000297, NT2RP2000420, NT2RP2001174,
 NT2RP2001233, NT2RP2001756, NT2RP2001869, NT2RP2002046, NT2RP2002252, NT2RP2002270,
 NT2RP2002464, NT2RP2002503, NT2RP2002520, NT2RP2002591, NT2RP2002880, NT2RP2002939,
 55 NT2RP2002993, NT2RP2003243, NT2RP2003329, NT2RP2003347, NT2RP2003480, NT2RP2003522,
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 NT2RP2004961, NT2RP2005003, NT2RP2005139, NT2RP2005325, NT2RP2005496, NT2RP2005701,
 NT2RP2005722, NT2RP2005776, NT2RP2005942, NT2RP2006238, NT2RP2006436, NT2RP3000050,

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	OVARC1002158,	OVARC1002165,	PLACE1000014,	PLACE1000213,	PLACE1000401,	PLACE1000562,
	PLACE1000611,	PLACE1000656,	PLACE1000712,	PLACE1 000793,	PLACE1000909,	PLACE1000948,
	PLACE1000977,	PLACE1001241,	PLACE1001257,	PLACE1001377,	PLACE1001517,	PLACE1001610,
5	PLACE1001761,	PLACE1001771,	PLACE1001817,	PLACE1001983,	PLACE1002046,	PLACE1002140,
	PLACE1002213,	PLACE1002395,	PLACE1002437,	PLACE1002500,	PLACE1002583,	PLACE1002714,
	PLACE1002722,	PLACE1002782,	PLACE1002794,	PLACE1002851,	PLACE1002908,	PLACE1003030,
	PLACE1003044,	PLACE1003045,	PLACE1003238,	PLACE1003296,	PLACE1003369,	PLACE1003420,
	PLACE1003493,	PLACE1003537,	PLACE1003553,	PLACE1003596,	PLACE1003760,	PLACE1003768,
10	PLACE1003771,	PLACE1003903,	PLACE1004149,	PLACE1004197,	PLACE1004203,	PLACE1004258,
	PLACE1004270,	PLACE1004277,	PLACE1004289,	PLACE1004473,	PLACE1004629,	PLACE1004646,
	PLACE1004743,	PLACE1004751,	PLACE1004793,	PLACE1004840,	PLACE1004969,	PLACE1005086,
	PLACE1005162,	PLACE1005206,	PLACE1005313,	PLACE1005467,	PLACE1005530,	PLACE1005595,
	PLACE1005611,	PLACE1005623,	PLACE1005763,	PLACE1005884,	PLACE1005890,	PLACE1005898,
15	PLACE1005934,	PLACE1005953,	PLACE1006157,	PLACE1006225,	PLACE1006239,	PLACE1006288,
	PLACE1006492,	PLACE1006534,	PLACE1006678,	PLACE1006754,	PLACE1006901,	PLACE1006935,
	PLACE1006956,	PLACE1007111,	PLACE1007243,	PLACE1007274,	PLACE1007282,	PLACE1007317,
	PLACE1007375,	PLACE1007386,	PLACE1007409,	PLACE1007416,	PLACE1007484,	PLACE1007583,
	PLACE1007632,	PLACE1007645,	PLACE1007649,	PLACE1007852,	PLACE1007877,	PLACE1007954,
20	PLACE1008273,	PLACE1008309,	PLACE1008331,	PLACE1008402,	PLACE1008424,	PLACE1008429,
	PLACE1008531,	PLACE1008532,	PLACE1008533,	PLACE1008568,	PLACE1008643,	PLACE1008693,
	PLACE1008715,	PLACE1009045,	PLACE1009094,	PLACE1009298,	PLACE1009319,	PLACE1009338,
	PLACE1009368,	PLACE1009493,	PLACE1009639,	PLACE1009659,	PLACE1009708,	PLACE1009731,
	PLACE1009845,	PLACE1009861,	PLACE1009935,	PLACE1009992,	PLACE1010089,	PLACE1010231,
25	PLACE1010321,	PLACE1010362,	PLACE1010599,	PLACE1010622,	PLACE1010662,	PLACE1010811,
	PLACE1010917,	PLACE1010942,	PLACE1010954,	PLACE1011090,	PLACE1011214,	PLACE1011221,
	PLACE1011371,	PLACE1011399,	PLACE1011492,	PLACE1011646,	PLACE1011749,	PLACE1011896,
	PLACE2000034,	PLACE2000062,	PLACE2000111,	PLACE2000132,	PLACE2000176,	PLACE2000187,
	PLACE2000216,	PLACE2000335,	PLACE2000341,	PLACE2000373,	PLACE2000379,	PLACE2000398,
30	PLACE2000399,	PLACE2000425,	PLACE2000438,	PLACE2000458,	PLACE2000477,	PLACE3000020,
	PLACE3000218,	PLACE3000226,	PLACE3000242,	PLACE3000244,	PLACE3000339,	PLACE3000373,
	PLACE3000399,	PLACE3000406,	PLACE3000413,	PLACE3000455,	PLACE4000052,	PLACE4000063,
	PLACE4000129,	PLACE4000247,	PLACE4000250,	PLACE4000259,	PLACE4000300,	PLACE4000387,
	PLACE4000431,	PLACE4000487,	PLACE4000494,	PLACE4000522,	PLACE4000548,	PLACE4000581,
35	PLACE4000593,	PLACE4000650,	THYRO1000156,	THYRO1000327,	THYRO1000394,	THYRO1000395,
	THYRO1000570,	THYRO1000748,	THYRO1000756,	THYRO1000783,	THYRO1001134,	THYRO1001271,
	THYRO1001287,	THYRO1001320,	THYRO1001401,	THYRO1001534,	THYRO1001537,	THYRO1001541,
	THYRO1001828,	Y79AA1000258,	Y79AA1000420,	Y79AA1000469,	Y79AA1000734,	Y79AA1000800,
	Y79AA1000976,	Y79AA1001023,	Y79AA1001177,	Y79AA1001384,	Y79AA1001394,	Y79AA1001603,
40	Y79AA1001647,	Y79AA1001846,	Y79AA1001874,	Y79AA1002139,	Y79AA1002246,	Y79AA1002351,
	Y79AA1002399,	Y79AA1002416,				
	[0276] The following 141 clones are categorized into glycoproteins-associated proteins.					
	HEMBA1000156,	HEMBA1000518,	HEMBA1000852,	HEMBA1001071,	HEMBA1001286,	HEMBA1001661,
	HEMBA1001734,	HEMBA1001866,	HEMBA1003071,	HEMBA1003077,	HEMBA1003281,	HEMBA1003538,
	HEMBA1003679,	HEMBA1003866,	HEMBA1005576,	HEMBA1005581,	HEMBA1005699,	HEMBA1006038,
45	HEMBA1006976,	HEMBA1007301,	HEMBA1000317,	HEMBA1000915,	HEMBA1001871,	HEMBA1001872,
	HEMBA1002193,	MAMMA1000672,	MAMMA1000897,	MAMMA1001030,	MAMMA1001388,	MAMMA1002329,
	MAMMA1002428,	MAMMA1002573,	MAMMA1003150,	NT2RM1000648,	NT2RM1001115,	NT2RM2000260,
	NT2RM2000407,	NT2RM2000422,	NT2RM2000490,	NT2RM2001499,	NT2RM2001659,	NT2RM2001930,
50	NT2RM4000820,	NT2RM4000857,	NT2RM4001810,	NT2RM4001813,	NT2RM4001987,	NT2RM4002145,
	NT2RM4002189,	NT2RM4002251,	NT2RM4002460,	NT2RM4002558,	NT2RP1000677,	NT2RP1000782,
	NT2RP1000856,	NT2RP1001546,	NT2RP2000056,	NT2RP2000070,	NT2RP2001295,	NT2RP2001378,
	NT2RP2001597,	NT2RP2001991,	NT2RP2002025,	NT2RP2002078,	NT2RP2002385,	NT2RP2004587,
	NT2RP2004732,	NT2RP2005531,	NT2RP3000207,	NT2RP3000531,	NT2RP3000825,	NT2RP3001140,
55	NT2RP3002810,	NT2RP3003672,	NT2RP3003701,	NT2RP3003716,	NT2RP3003914,	NT2RP3004148,
	NT2RP4000212,	NT2RP4000417,	NT2RP4000724,	NT2RP4000817,	NT2RP4000925,	NT2RP4001150,
	NT2RP4001372,	NT2RP4001730,	NT2RP4001822,	NT2RP4001823,	NT2RP5003522,	OVARC1000091,
	OVARC1000288,	OVARC1000682,	OVARC1001055,	OVARC1001506,	OVARC1001713,	OVARC1002127,
	PLACE1000213,	PLACE1000401,	PLACE1002437,	PLACE1002583,		

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	NT2RM2000581,	NT2RM2000609,	NT2RM2000821,	NT2RM2001370,	NT2RM2001393,	NT2RM2001499,
	NT2RM2001547,	NT2RM2001613,	NT2RM2001648,	NT2RM2001659,	NT2RM2001671,	NT2RM2001688,
	NT2RM2001698,	NT2RM2001718,	NT2RM2001753,	NT2RM2001760,	NT2RM2001785,	NT2RM2001930,
	NT2RM2001950,	NT2RM2001997,	NT2RM2001998,	NT2RM2002049,	NT2RM2002145,	NT2RM4000233,
5	NT2RM4000433,	NT2RM4000457,	NT2RM4000486,	NT2RM4000496,	NT2RM4000520,	NT2RM4000634,
	NT2RM4000674,	NT2RM4000700,	NT2RM4000764,	NT2RM4000778,	NT2RM4000795,	NT2RM4000820,
	NT2RM4000857,	NT2RM4001032,	NT2RM4001054,	NT2RM4001116,	NT2RM4001455,	NT2RM4001666,
	NT2RM4001810,	NT2RM4001813,	NT2RM4001930,	NT2RM4001987,	NT2RM4002054,	NT2RM4002073,
	NT2RM4002145,	NT2RM4002146,	NT2RM4002189,	NT2RM4002194,	NT2RM4002251,	NT2RM4002339,
10	NT2RM4002438,	NT2RM4002446,	NT2RM4002452,	NT2RM4002460,	NT2RM4002493,	NT2RM4002558,
	NT2RM4002565,	NT2RM4002571,	NT2RM4002594,	NT2RP1000130,	NT2RP1000191,	NT2RP1000326,
	NT2RP1000358,	NT2RP1000413,	NT2RP1000418,	NT2RP1000547,	NT2RP1000609,	NT2RP1000677,
	NT2RP1000767,	NT2RP1000782,	NT2RP1000856,	NT2RP1001113,	NT2RP1001247,	NT2RP1001286,
	NT2RP1001310,	NT2RP1001311,	NT2RP1001313,	NT2RP1001385,	NT2RP1001449,	NT2RP1001546,
15	NT2RP1001569,	NT2RP2000032,	NT2RP2000040,	NT2RP2000056,	NT2RP2000070,	NT2RP2000091,
	NT2RP2000114,	NT2RP2000120,	NT2RP2000173,	NT2RP2000175,	NT2RP2000195,	NT2RP2000257,
	NT2RP2000270,	NT2RP2000283,	NT2RP2000288,	NT2RP2000289,	NT2RP2000459,	NT2RP2000516,
	NT2RP2000660,	NT2RP2000842,	NT2RP2000892,	NT2RP2001081,	NT2RP2001268,	NT2RP2001295,
	NT2RP2001366,	NT2RP2001378,	NT2RP2001576,	NT2RP2001581,	NT2RP2001597,	NT2RP2001613,
20	NT2RP2001947,	NT2RP2001991,	NT2RP2002025,	NT2RP2002066,	NT2RP2002078,	NT2RP2002105,
	NT2RP2002312,	NT2RP2002325,	NT2RP2002385,	NT2RP2002479,	NT2RP2002537,	NT2RP2002643,
	NT2RP2002701,	NT2RP2002740,	NT2RP2002857,	NT2RP2003125,	NT2RP2003297,	NT2RP2003433,
	NT2RP2003446,	NT2RP2003466,	NT2RP2003506,	NT2RP2003513,	NT2RP2003629,	NT2RP2003668,
	NT2RP2003760,	NT2RP2003777,	NT2RP2003781,	NT2RP2004041,	NT2RP2004142,	NT2RP2004194,
25	NT2RP2004270,	NT2RP2004300,	NT2RP2004392,	NT2RP2004655,	NT2RP2004681,	NT2RP2004775,
	NT2RP2004799,	NT2RP2004936,	NT2RP2004959,	NT2RP2005012,	NT2RP2005159,	NT2RP2005227,
	NT2RP2005270,	NT2RP2005344,	NT2RP2005465,	NT2RP2005509,	NT2RP2005752,	NT2RP2005781,
	NT2RP2005784,	NT2RP2005812,	NT2RP2006069,	NT2RP2006100,	NT2RP2006141,	NT2RP2006184,
	NT2RP2006261,	NT2RP2006565,	NT2RP2006571,	NT2RP2006573,	NT2RP3000092,	NT2RP3000109,
30	NT2RP3000134,	NT2RP3000207,	NT2RP3000333,	NT2RP3000341,	NT2RP3000393,	NT2RP3000439,
	NT2RP3000441,	NT2RP3000531,	NT2RP3000685,	NT2RP3000825,	NT2RP3000826,	NT2RP3000852,
	NT2RP3000919,	NT2RP3001084,	NT2RP3001096,	NT2RP3001126,	NT2RP3001140,	NT2RP3001176,
	NT2RP3001260,	NT2RP3001282,	NT2RP3001355,	NT2RP3001383,	NT2RP3001426,	NT2RP3001453,
	NT2RP3001497,	NT2RP3001538,	NT2RP3001589,	NT2RP3001642,	NT2RP3001708,	NT2RP3001716,
35	NT2RP3001727,	NT2RP3001739,	NT2RP3001799,	NT2RP3001943,	NT2RP3001944,	NT2RP3002002,
	NT2RP3002007,	NT2RP3002014,	NT2RP3002054,	NT2RP3002108,	NT2RP3002163,	NT2RP3002351,
	NT2RP3002455,	NT2RP3002549,	NT2RP3002602,	NT2RP3002628,	NT2RP3002650,	NT2RP3002687,
	NT2RP3002701,	NT2RP3002810,	NT2RP3002869,	NT2RP3002969,	NT2RP3002985,	NT2RP3003008,
	NT2RP3003059,	NT2RP3003071,	NT2RP3003101,	NT2RP3003145,	NT2RP3003197,	NT2RP3003203,
40	NT2RP3003242,	NT2RP3003302,	NT2RP3003353,	NT2RP3003409,	NT2RP3003576,	NT2RP3003621,
	NT2RP3003665,	NT2RP3003672,	NT2RP3003701,	NT2RP3003716,	NT2RP3003799,	NT2RP3003828,
	NT2RP3003914,	NT2RP3003918,	NT2RP3003992,	NT2RP3004051,	NT2RP3004148,	NT2RP3004155,
	NT2RP3004207,	NT2RP3004282,	NT2RP3004454,	NT2RP3004480,	NT2RP3004503,	NT2RP4000008,
	NT2RP4000051,	NT2RP4000151,	NT2RP4000212,	NT2RP4000243,	NT2RP4000259,	NT2RP4000323,
45	NT2RP4000417,	NT2RP4000500,	NT2RP4000524,	NT2RP4000556,	NT2RP4000560,	NT2RP4000588,
	NT2RP4000713,	NT2RP4000724,	NT2RP4000817,	NT2RP4000833,	NT2RP4000878,	NT2RP4000907,
	NT2RP4000925,	NT2RP4000928,	NT2RP4000973,	NT2RP4000989,	NT2RP4001057,	NT2RP4001064,
	NT2RP4001079,	NT2RP4001117,	NT2RP4001138,	NT2RP4001149,	NT2RP4001150,	NT2RP4001174,
	NT2RP4001219,	NT2RP4001274,	NT2RP4001313,	NT2RP4001345,	NT2RP4001372,	NT2RP4001373,
50	NT2RP4001379,	NT2RP4001498,	NT2RP4001547,	NT2RP4001571,	NT2RP4001574,	NT2RP4001644,
	NT2RP4001656,	NT2RP4001677,	NT2RP4001730,	NT2RP4001739,	NT2RP4001803,	NT2RP4001822,
	NT2RP4001823,	NT2RP4001950,	NT2RP4001975,	NT2RP4002052,	NT2RP4002075,	NT2RP5003500,
	NT2RP5003506,	NT2RP5003522,	NT2RP5003534,	OVARC1000060,	OVARC1000335,	OVARC1000682,
	OVARC1000689,	OVARC1000700,	OVARC1000722,	OVARC1000751,	OVARC1000850,	OVARC1000890,
55	OVARC1000924,	OVARC1000936,	OVARC1000959,	OVARC1000984,	OVARC1000999,	OVARC1001034,
	OVARC1001055,	OVARC1001117,	OVARC1001129,	OVARC1001154,	OVARC1001329,	OVARC1001381,
	OVARC1001391,	OVARC1001453,	OVARC1001476,	OVARC1001506,	OVARC1001610,	OVARC1001702,
	OVARC1001703,	OVARC1001713,	OVARC1001745,	OVARC1001767,	OVARC1002127,	OVARC1002138,

[0267] The clones that presumably encode proteins belonging to the category of DNA- and/or RNA-binding proteins are those which matched the data with the terms "DNA-binding" or "RNA-binding".

[0268] The clones that presumably encode proteins belonging to the category of RNA synthesis-associated proteins are those which matched the data with the terms "RNA splicing", "RNA processing", "RNA helicase", or "polyadenylation".

[0269] The clones that presumably encode proteins belonging to the category of protein synthesis- and/or protein transport-associated proteins are those which matched the data with the terms "translation regulation", "protein biosynthesis", "amino-acid biosynthesis", "ribosomal protein", "protein transport", or "signal recognition particle".

[0270] The clones that presumably encode proteins belonging to the category of cytoskeleton-associated proteins are those which matched the data with the terms "structural protein", "cytoskeleton", "actin-binding", or "microtubules".

[0271] The clones that presumably encode proteins belonging to the category of cell division- and/or cell proliferation-associated proteins are those which matched the data with the terms "cell division", "cell cycle", "mitosis", "chromosomal protein", "cell growth", or "apoptosis".

[0272] The clones that presumably encode proteins belonging to the category of embryogenesis- and/or development-associated proteins are those which matched the data with the terms "developmental protein".

[0273] The clones that presumably encode proteins belonging to the category of cellular defense-associated proteins are those which matched the data with the terms "heat shock", "DNA repair", or "DNA damage".

[0274] When a clone belonged to the above-mentioned multiple functional categories, the clone was classified into the multiple categories. However, the functions of the protein encoded by the clone are not limited to the functions of the categories into which the clone was classified, and therefore, additional functions can be found for the protein by further analyses.

[0275] The following 796 clones are categorized into secretory or membrane proteins.

	HEMBA1000356,	HEMBA1000518,	HEMBA1000531,	HEMBA1000637,	HEMBA1000719,	HEMBA1000817,
	HEMBA1000822,	HEMBA1000852,	HEMBA1000870,	HEMBA1000991,	HEMBA1001052,	HEMBA1001071,
25	HEMBA1001085,	HEMBA1001286,	HEMBA1001351,	HEMBA1001407,	HEMBA1001446,	HEMBA1001515,
	HEMBA1001557,	HEMBA1001569,	HEMBA1001661,	HEMBA1001734,	HEMBA1001746,	HEMBA1001866,
	HEMBA1002125,	HEMBA1002150,	HEMBA1002166,	HEMBA1002417,	HEMBA1002462,	HEMBA1002475,
	HEMBA1002477,	HEMBA1002486,	HEMBA1002609,	HEMBA1002659,	HEMBA1002661,	HEMBA1002780,
	HEMBA1002818,	HEMBA1002876,	HEMBA1002921,	HEMBA1003071,	HEMBA1003077,	HEMBA1003079,
30	HEMBA1003086,	HEMBA1003096,	HEMBA1003281,	HEMBA1003286,	HEMBA1003538,	HEMBA1003711,
	HEMBA1003742,	HEMBA1003803,	HEMBA1004055,	HEMBA1004143,	HEMBA1004146,	HEMBA1004207,
	HEMBA1004341,	HEMBA1004461,	HEMBA1004577,	HEMBA1004637,	HEMBA1004752,	HEMBA1004756,
	HEMBA1004850,	HEMBA1004889,	HEMBA1004923,	HEMBA1004930,	HEMBA1005029,	HEMBA1005035,
	HEMBA1005050,	HEMBA1005552,	HEMBA1005576,	HEMBA1005581,	HEMBA1005588,	HEMBA1005616,
35	HEMBA1005699,	HEMBA1005991,	HEMBA1006036,	HEMBA1006038,	HEMBA1006067,	HEMBA1006173,
	HEMBA1006198,	HEMBA1006293,	HEMBA1006310,	HEMBA1006492,	HEMBA1006502,	HEMBA1006583,
	HEMBA1006659,	HEMBA1006758,	HEMBA1006789,	HEMBA1006921,	HEMBA1006926,	HEMBA1006976,
	HEMBA1007203,	HEMBA1007301,	HEMBA1000037,	HEMBA1000050,	HEMBA1000054,	HEMBA1000175,
	HEMBA1000317,	HEMBA1000556,	HEMBA1000593,	HEMBA1000631,	HEMBA1000763,	HEMBA1000827,
40	HEMBA1000915,	HEMBA1000975,	HEMBA1001112,	HEMBA1001151,	HEMBA1001177,	HEMBA1001302,
	HEMBA1001348,	HEMBA1001564,	HEMBA1001630,	HEMBA1001871,	HEMBA1001872,	HEMBA1001925,
	HEMBA1001962,	HEMBA1002042,	HEMBA1002044,	HEMBA1002142,	HEMBA1002190,	HEMBA1002193,
	HEMBA1002247,	HEMBA1002383,	HEMBA1002387,	HEMBA1002550,	HEMBA1002600,	HEMBA1002692,
	MAMMA1000045,	MAMMA1000129,	MAMMA1000133,	MAMMA1000277,	MAMMA1000278,	MAMMA1000410,
45	MAMMA1000416,	MAMMA1000472,	MAMMA1000672,	MAMMA1000684,	MAMMA1000714,	MAMMA1000734,
	MAMMA1000778,	MAMMA1000798,	MAMMA1000842,	MAMMA1000859,	MAMMA1000897,	MAMMA1000956,
	MAMMA1001008,	MAMMA1001030,	MAMMA1001041,	MAMMA1001073,	MAMMA1001080,	MAMMA1001139,
	MAMMA1001154,	MAMMA1001322,	MAMMA1001388,	MAMMA1001411,	MAMMA1001487,	MAMMA1001751,
	MAMMA1001754,	MAMMA1001771,	MAMMA1002009,	MAMMA1002427,	MAMMA1002428,	MAMMA1002461,
50	MAMMA1002524,	MAMMA1002573,	MAMMA1002598,	MAMMA1002655,	MAMMA1002684,	MAMMA1002769,
	MAMMA1002844,	MAMMA1002881,	MAMMA1002890,	MAMMA1002938,	MAMMA1002947,	MAMMA1003035,
	MAMMA1003089,	MAMMA1003146,	MAMMA1003150,	NT2RM1000035,	NT2RM1000037,	NT2RM1000062,
	NT2RM1000080,	NT2RM1000092,	NT2RM1000131,	NT2RM1000199,	NT2RM1000257,	NT2RM1000260,
	NT2RM1000355,	NT2RM1000430,	NT2RM1000563,	NT2RM1000648,	NT2RM1000742,	NT2RM1000770,
55	NT2RM1000800,	NT2RM1000811,	NT2RM1000833,	NT2RM1000857,	NT2RM1000867,	NT2RM1000882,
	NT2RM1000905,	NT2RM1001008,				
	NT2RM1001115,	NT2RM1001139,	NT2RM2000259,	NT2RM2000260,	NT2RM2000287,	NT2RM2000395,
	NT2RM2000402,	NT2RM2000407,	NT2RM2000422,	NT2RM2000490,	NT2RM2000522,	NT2RM2000566,

Y79AA1000752, Y79AA1000774, Y79AA1000782, Y79AA1000784, Y79AA1000794, Y79AA1000800,
 Y79AA1000802, Y79AA1000805, Y79AA1000824, Y79AA1000827, Y79AA1000833, Y79AA1000850,
 Y79AA1000962, Y79AA1000966, Y79AA1000968, Y79AA1000969, Y79AA1000976, Y79AA1000985,
 Y79AA1001023, Y79AA1001041, Y79AA1001048, Y79AA1001061, Y79AA1001068, Y79AA1001077,
 5 Y79AA1001078, Y79AA1001145, Y79AA1001167, Y79AA1001177, Y79AA1001185, Y79AA1001211,
 Y79AA1001216, Y79AA1001228, Y79AA1001233, Y79AA1001236, Y79AA1001281, Y79AA1001299,
 Y79AA1001312, Y79AA1001323, Y79AA1001384, Y79AA1001391, Y79AA1001394, Y79AA1001402,
 Y79AA1001493, Y79AA1001511, Y79AA1001533, Y79AA1001541, Y79AA1001548, Y79AA1001555,
 Y79AA1001581, Y79AA1001585, Y79AA1001594, Y79AA1001603, Y79AA1001613, Y79AA1001647,
 10 Y79AA1001665, Y79AA1001679, Y79AA1001692, Y79AA1001696, Y79AA1001705, Y79AA1001711,
 Y79AA1001781, Y79AA1001805, Y79AA1001827, Y79AA1001846, Y79AA1001866, Y79AA1001874,
 Y79AA1001875, Y79AA1001923, Y79AA1001963, Y79AA1002027, Y79AA1002083, Y79AA1002089,
 Y79AA1002103, Y79AA1002115, Y79AA1002125, Y79AA1002139, Y79AA1002204, Y79AA1002208,
 Y79AA1002209, Y79AA1002210, Y79AA1002211, Y79AA1002220, Y79AA1002229, Y79AA1002234,
 15 Y79AA1002246, Y79AA1002258, Y79AA1002298, Y79AA1002307, Y79AA1002311, Y79AA1002351,
 Y79AA1002361, Y79AA1002399, Y79AA1002407, Y79AA1002416, Y79AA1002431, Y79AA1002433,
 Y79AA1002472, Y79AA1002482, Y79AA1002487,

[0258] Among the 4997 clones, there are 2189 clones that presumably encode proteins belonging to any of the categories of secretory or membrane proteins, glycoprotein-associated proteins, signal transduction-associated proteins, transcription-associated proteins, disease-associated proteins, enzymes and/or metabolism-associated proteins, ATP- and/or GTP-binding proteins, nuclear proteins, DNA- and/or RNA-binding proteins, RNA synthesis-associated proteins, protein synthesis- and/or protein transport-associated proteins, cytoskeleton-associated proteins, cell division- and/or cell proliferation-associated proteins, embryogenesis- and/or development-associated proteins, or cellular defense-associated proteins.

[0259] The clones that presumably encode proteins belonging to the category of secretory or membrane proteins are those which matched the full-length sequences of Swiss-Prot database with the keywords "growth factor", "cytokine", "hormone", "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", or "connective tissue"; those which matched the data, suggesting that the proteins are secretory or membrane proteins; or those which matched the full-length sequences of GenBank or UniGene database with similar description; and, further, those predicted to have an N-terminal signal sequence or a transmembrane region as a result of domain search for the amino acid sequences deduced from the full-length nucleotide sequences.

[0260] The clones that presumably encode proteins belonging to the category of glycoprotein-associated proteins are those which matched the full-length sequences of Swiss-Prot database with the keywords "glycoprotein"; those which matched the data, suggesting that the proteins are glycoprotein; or those which matched the full-length sequences of GenBank or UniGene database with similar description.

[0261] The clones that presumably encode proteins belonging to the category of signal transduction-associated proteins are those which matched the full-length sequences of Swiss-Prot database with the keywords "serine/threonine-protein kinase", "tyrosine-protein kinase", or "SH3 domain"; those which matched the data, suggesting that the proteins are signal transduction-associated proteins (for example, "ADP-ribosylation factor"); or those which matched the full-length sequences of GenBank or UniGene database with similar description.

[0262] The clones that presumably encode proteins belonging to the category of transcription-associated proteins are those which matched the full-length sequences of Swiss-Prot database with the keywords "transcription regulation", "zinc finger", or "homeobox"; those which matched the data, suggesting that the proteins are transcription-associated proteins; or those which matched the full-length sequences of GenBank or UniGene database with similar description.

[0263] The clones that presumably encode proteins belonging to the category of disease-associated proteins are those which matched the full-length sequences of Swiss-Prot database with the keywords "disease mutation" or "syndrome"; those which matched the data, suggesting that the proteins are disease-associated proteins; or those which matched the full-length sequences of Swiss-Prot database and GenBank or UniGene database where the matched sequences of genes or proteins which had been registered in the database of Online Mendelian Inheritance in Man (OMIM) (<http://www.ncbi.nlm.nih.gov/Omim/>), which is a database of human genes and diseases.

[0264] The clones that presumably encode proteins belonging to the category of enzymes and/or metabolism-associated proteins are those which showed the terms "metabolism", "oxidoreductase", or "E.C. No. (Enzyme commission number)" in the matching data.

[0265] The clones that presumably encode proteins belonging to the category of ATP- and/or GTP-binding proteins are those which matched the data with the terms "ATP-binding" or "GTP-binding".

[0266] The clones that presumably encode proteins belonging to the category of nuclear proteins are those which matched the data with the terms "nuclear protein".

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Y79AA1000342//Zinc finger, C2H2 type
 Y79AA1000349//Double-stranded RNA binding motif
 Y79AA1000627//Zinc finger, C2H2 type
 Y79AA1000705//Helicases conserved C-terminal domain
 5 Y79AA1000752//KH domain family of RNA binding proteins
 Y79AA1000833//Tubulin
 Y79AA1001048//Acyl-CoA dehydrogenases
 Y79AA1001391//Homeobox domain
 Y79AA1001394//ATPases associated with various cellular activities (AAA)
 10 Y79AA1001493//Ubiquitin-conjugating enzymes
 Y79AA1001613//Zinc finger, C2H2 type
 Y79AA1001874//TNFR/NGFR cysteine-rich region
 Y79AA1002027//Ubiquitin-conjugating enzymes
 Y79AA1002139//DnaJ, prokaryotic heat shock protein
 15 Y79AA1002208//Ank repeat
 Y79AA1002246//C2 domain
 Y79AA1002307//Fibronectin type III domain
 Y79AA1002472//Zinc finger, C2H2 type
 HEMBA1003538//CUB domain HEMBA1003645//WD domain, G-beta repeats //Src homology domain 3
 20 HEMBA1005206//Glutathione S-transferases.
 HEMBA1006521//Alcohol/other dehydrogenases, short chain type
 HEMBB1001482//Zinc finger, C2H2 type HEMBB1001915//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2 HEMBB1002044//Cadherin MAMMA1000183//Zinc finger, C2H2 type
 MAMMA1000897//von Willebrand factor type A domain MAMMA1001080//IG superfamily MAMMA1002498//IG superfamily MAMMA1002573//KH domain family of RNA binding proteins MAMMA1002617//Zinc finger, C2H2 type
 25 NT2RM1000833//eubacterial secY protein NT2RM2001797//Zinc finger, C2H2 type
 NT2RP1001013//Zinc finger, C2H2 type NT2RP2001233//Zinc finger, C2H2 type
 NT2RP2001440//14-3-3 proteins NT2RP2002105//7 transmembrane receptor (rhodopsin family)
 NT2RP3001723//Laminin G domain NT2RP3001938//Eukaryotic protein kinase domain NT2RP3002330//Elongation factor Tu family (contains ATP/GTP binding P-loop) NT2RP3003133//Zinc finger, C2H2 type
 30 NT2RP3003500//Eukaryotic protein kinase domain NT2RP3003799//C2 domain
 NT2RP3003800//Eukaryotic protein kinase domain NT2RP3004013//Double-stranded RNA binding motif
 NT2RP3004125//Zinc finger, C2H2 type
 OVARC1001244//Bromodomain OVARC1001496//D-isomer specific 2-hydroxyacid dehydrogenases
 35 PLACE1000007//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 PLACE1001118//Zinc finger, C2H2 type PLACE1010310//Zinc finger, C2H2 type PLACE1011896//wnt family of developmental signaling proteins PLACE3000124//Src homology domain 2
 PLACE4000100//D-isomer specific 2-hydroxyacid dehydrogenases
 PLACE4000259//Helicases conserved C-terminal domain PLACE4000261//Bromodomain SKNMC1000013//ABC transporters SKNMC1000091//Basic region plus leucine zipper transcription factors THYRO1000343//Src homology domain 3 THYRO1000569//Zinc finger, C2H2 type THYRO1001189//Zinc finger, C2H2 type Y79AA1002103//Zinc finger, C2H2 type PLACE3000350//Eukaryotic protein kinase domain
 40 PLACE4000156//Zinc finger, C2H2 type

45 EXAMPLE 18

Classification of cDNA clones into functional categories based on the full-length nucleotide sequences

50 **[0257]** Prediction of functions of proteins encoded by the clones and the categorization thereof were performed based on the results of homology search (see Homology search results 6, 12, 13 and 14) of the databases, GenBank, Swiss-Prot and UniGene, for the full-length nucleotide sequences of 4997 clones and based on the results of domain search (see Example 17) of the deduced amino acid sequences encoded by the full-length nucleotide sequences. The target 4997 clones are listed below:

55 HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000030, HEMBA1000042, HEMBA1000046,
 HEMBA1000050, HEMBA1000076, HEMBA1000129, HEMBA1000141, HEMBA1000150, HEMBA1000156,
 HEMBA1000158, HEMBA1000168, HEMBA1000185, HEMBA1000193, HEMBA1000201, HEMBA1000213,
 HEMBA1000216, HEMBA1000227, HEMBA1000231, HEMBA1000243, HEMBA1000244, HEMBA1000251,
 HEMBA1000264, HEMBA1000280, HEMBA1000282, HEMBA1000288, HEMBA1000290, HEMBA1000302,

PLACE1009099//Zinc finger, C2H2 type
 PLACE1009246//LIM domain containing proteins
 PLACE1009468//WD domain, G-beta repeats
 PLACE1009476//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 5 PLACE1009524//PH (pleckstrin homology) domain
 PLACE1009596//WD domain, G-beta repeats
 PLACE1009622//Double-stranded RNA binding motif
 PLACE1009861//Cysteine proteases
 PLACE1009925//Helicases conserved C-terminal domain
 10 PLACE1009992//CUB domain //EGF-like domain //Sushi domain //Trypsin
 PLACE1010053//Double-stranded RNA binding motif
 PLACE1010089//Ubiquitin carboxyl-terminal hydrolases family 2
 PLACE1010702//Zinc finger, C2H2 type
 PLACE1010833//EF hand
 15 PLACE1010926//Src homology domain 3
 PLACE1010960//Actins
 PLACE1011041//Src homology domain 3
 PLACE1011046//C2 domain //Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-spe-
 cific phospholipase C, Y domain
 20 PLACE1011114//Helicases conserved C-terminal domain
 PLACE1011160//Zinc finger, C3HC4 type (RING finger)
 PLACE1011263//Ank repeat
 PLACE1011433//Zinc finger, C2H2 type
 PLACE1011576//Zinc finger, C2H2 type
 25 PLACE1011923//Eukaryotic protein kinase domain
 PLACE2000034//Fibronectin type III domain //IG superfamily
 PLACE2000072//Zinc finger, C2H2 type
 PLACE2000111//IG superfamily
 PLACE2000164//WD domain, G-beta repeats
 30 PLACE2000216//PH (pleckstrin homology) domain
 PLACE2000341//Sodium:solute symporter family
 PLACE2000371//Src homology domain 2
 PLACE2000373//Thrombospondin type 1 domain
 PLACE2000398//IG superfamily
 35 PLACE2000427//Helicases conserved C-terminal domain
 PLACE2000458//Cadherin
 PLACE3000020//Guanylate cyclases
 PLACE3000169//Zinc finger, C2H2 type
 PLACE4000014//Helicases conserved C-terminal domain
 40 PLACE4000052//ABC transporters
 PLACE4000192//Zinc finger, C2H2 type
 PLACE4000211//Bromodomain
 PLACE4000431//Helicases conserved C-terminal domain
 PLACE4000522//Ank repeat
 45 PLACE4000581//EGF-like domain //Sushi domain
 PLACE4000654//Ubiquitin-conjugating enzymes
 THYRO1000072//IG superfamily
 THYRO1000242//Zinc finger, C2H2 type
 THYRO1000288//Zinc-binding metalloprotease domain
 50 THYRO1000488//Zinc finger, C3HC4 type (RING finger)
 THYRO1000501//Zinc finger, C3HC4 type (RING finger)
 THYRO1000666//Kinesin motor domain
 THYRO1000748//Src homology domain 3
 THYRO1000926//3' 5'-cyclic nucleotide phosphodiesterases
 55 THYRO1001661//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 THYRO1001671//Ubiquitin family
 Y79AA1000037//Zinc finger, C3HC4 type (RING finger)
 Y79AA1000214//Core histones H2A, H2B, H3 and H4

PLACE1002571//Actins
 PLACE1002685//Src homology domain 2
 PLACE1002722//7 transmembrane receptor (rhodopsin family)
 PLACE1002775//Bromodomain
 5 PLACE1002834//Zinc finger, C2H2 type
 PLACE1003100//Alcohol/other dehydrogenases, short chain type
 PLACE1003174//Ubiquitin-conjugating enzymes
 PLACE1003238//7 transmembrane receptor (rhodopsin family)
 PLACE1003302//Zinc finger, C2H2 type
 10 PLACE1003334//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1003366//C2 domain
 PLACE1003394//Ras family (contains ATP/GTP binding P-loop)
 PLACE1003420//Mitochondrial carrier proteins
 PLACE1003493//C1q domain
 15 PLACE1003519//KH domain family of RNA binding-proteins
 PLACE1003723//Src homology domain 2
 PLACE1003738//Zinc finger, C2H2 type
 PLACE1003888//C2 domain //Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 20 PLACE1004128//WD domain, G-beta repeats
 PLACE1004358//PH (pleckstrin homology) domain
 PLACE1004428//Acyl-CoA dehydrogenases
 PLACE1004437//Isocitrate and isopropylmalate dehydrogenases
 PLACE1004506//LIM domain containing proteins
 25 PLACE1004674//EF hand
 PLACE1004918//L-lactate dehydrogenases
 PLACE1005243//Eukaryotic protein kinase domain
 PLACE1005305//Adenylate kinases
 PLACE1005327//Src homology domain 3
 30 PLACE1005530//Zinc finger, C3HC4 type (RING finger)
 PLACE1005646//Helicases conserved C-terminal domain
 PLACE1005656//Ribonucleotide reductases
 PLACE1005966//WD domain, G-beta repeats
 PLACE1006157//Sushi domain
 35 PLACE1006196//DEAH and DEAR box helicases //Helicases conserved C-terminal domain
 PLACE1006438//Zinc finger, C2H2 type
 PLACE1006626//Double-stranded RNA binding motif
 PLACE1006754//IG superfamily
 PLACE1006829//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 40 PLACE1006917//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1006956//ABC transporters
 PLACE1006958//Heat shock hsp70 proteins
 PLACE1007375//C2 domain
 PLACE1007488//PH (pleckstrin homology) domain
 45 PLACE1007511//Intermediate filament proteins
 PLACE1007537//Ank repeat
 PLACE1007544//Zinc finger, C2H2 type
 PLACE1007547//Zinc finger, C3HC4 type (RING finger)
 PLACE1007598//Zinc finger, C2H2 type
 50 PLACE1007697//ABC transporters
 PLACE1007958//3'5'-cyclic nucleotide phosphodiesterases
 PLACE1007969//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1008201//Zinc finger, C2H2 type
 PLACE1008429//Ank repeat
 55 PLACE1008465//Zinc finger, C2H2 type
 PLACE1008650//WD domain, G-beta repeats
 PLACE1009020//Aminotransferases class-V
 PLACE1009094//von Willebrand factor type C domain

NT2RP4001822//4 transmembrane segments integral membrane proteins
 NT2RP4001823//Fibrinogen beta and gamma chains, C-terminal globular domain
 NT2RP4001893//Ank repeat
 NT2RP4001896//WD domain, G-beta repeats
 5 NT2RP4001927//WD domain, G-beta repeats
 NT2RP4001938//Zinc finger, C2H2 type
 NT2RP4002047//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP4002078//Zinc finger, C2H2 type
 NT2RP4002408//Eukaryotic protein kinase domain
 10 NT2RP4002905//Cyclins
 NT2RP5003477//WD domain, G-beta repeats
 OVARC1000006//Core histones H2A, H2B, H3 and H4
 OVARC1000085//Proteasome A-type and B-type
 OVARC1000148//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 15 OVARC1000556//Eukaryotic protein kinase domain
 OVARC1000649//PH (pleckstrin homology) domain //Src homology domain 2
 OVARC1000746//Double-stranded RNA binding motif
 OVARC1000885//Alcohol/other dehydrogenases, short chain type
 OVARC1000937//Cyclins
 20 OVARC1000999//Ank repeat
 OVARC1001154//Granulins
 OVARC1001180//Ubiquitin family
 OVARC1001306//Helix-loop-helix DNA-binding domain
 OVARC1001577//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 25 OVARC1001731//Tropomyosins
 OVARC1001943//Zinc finger, C2H2 type
 OVARC1002050//Spectrin alpha chain, repeated domain
 OVARC1002112//Core histones H2A, H2B, H3 and H4
 OVARC1002138//ATPases associated with various cellular activities (AAA)
 30 OVARC1002182//WD domain, G-beta repeats
 PLACE1000014//Zinc finger, C3HC4 type (RING finger)
 PLACE1000040//Ras family (contains ATP/GTP binding P-loop)
 PLACE1000050//Zinc finger, C2H2 type
 PLACE1000081//PH (pleckstrin homology) domain
 35 PLACE1000142//Enoyl-CoA hydratase/isomerase
 PLACE1000401//IG superfamily
 PLACE1000406//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 PLACE1000420//Bacterial mutT protein
 PLACE1000706//Bromodomain
 40 PLACE1000769//KH domain family of RNA binding proteins
 PLACE1000786//PH (pleckstrin homology) domain
 PLACE1000863//Ribosomal protein S4
 PLACE1000909//Ank repeat
 PLACE1000972//Src homology domain 3
 45 PLACE1000979//Zinc finger, C2H2 type
 PLACE1001304//Zinc finger, C2H2 type
 PLACE1001387//Src homology domain 3
 PLACE1001632//Zinc finger, C2H2 type
 PLACE1001672//Aminotransferases class-III pyridoxal-phosphate
 50 PLACE1001716//Zinc finger, CCHC class
 PLACE1001739//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 PLACE1001781//Phosphoglucomutase and phosphomannomutase phosphoserine
 PLACE1001869//FGGY family of carbohydrate kinases
 PLACE1002438//Zinc finger, C2H2 type
 55 PLACE1002450//Zinc finger, C2H2 type
 PLACE1002474//EGF-like domain //von Willebrand factor type A domain
 PLACE1002499//Zinc finger, C3HC4 type (RING finger)
 PLACE1002532//Homeobox domain

NT2RP3002628//DnaJ, prokaryotic heat shock protein //Thioredoxins
 NT2RP3002663//PH (pleckstrin homology) domain
 NT2RP3002909//Ank repeat
 NT2RP3002953//Cadherin
 5 NT2RP3002969//AMP-binding enzymes
 NT2RP3003061//Ank repeat
 NT2RP3003145//Zinc carboxypeptidases
 NT2RP3003230//WD domain, G-beta repeats
 NT2RP3003251//Zinc finger, C3HC4 type (RING finger)
 10 NT2RP3003278//Ank repeat //Zinc finger, C2H2 type
 NT2RP3003282//PH (pleckstrin homology) domain
 NT2RP3003311//PH (pleckstrin homology) domain
 NT2RP3003385//Ank repeat //Chaperonins clpA/B
 NT2RP3003589//Ras family (contains ATP/GTP binding P-loop)
 15 NT2RP3003621//CUB domain //Kringle domain
 NT2RP3003701//Thrombospondin type 1 domain
 NT2RP3003716//Fibronectin type III domain
 NT2RP3003809//ATPases associated with various cellular activities (AAA)
 NT2RP3004016//Zinc finger, C3HC4 type (RING finger)
 20 NT2RP3004207//CUB domain //Sushi domain
 NT2RP3004209//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP3004242//PH (pleckstrin homology) domain
 NT2RP3004262//DnaJ, prokaryotic heat shock protein
 NT2RP3004566//Zinc finger, C2H2 type
 25 NT2RP3004569//Ank repeat
 NT2RP3004594//HMG (high mobility group) box
 NT2RP3004617//Zinc finger, C3HC4 type (RING finger)
 NT2RP4000259//Glutathione peroxidases
 NT2RP4000370//Prokaryotic-type class I peptide chain release factors
 30 NT2RP4000376//WD domain, G-beta repeats
 NT2RP4000398//Zinc finger, C2H2 type
 NT2RP4000455//Forkhead-associated (FHA) domain //Zinc finger, C3HC4 type (RING finger)
 NT2RP4000457//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP4000518//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 35 NT2RP4000588//Actinin-type actin-binding domain containing proteins
 NT2RP4000614//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP4000648//Forkhead-associated (FHA) domain //Zinc finger, C3HC4 type (RING finger)
 NT2RP4000837//Zinc finger, C2H2 type
 NT2RP4000839//WD domain, G-beta repeats
 40 NT2RP4000865//Zinc finger, C2H2 type
 NT2RP4000907//Fibronectin type III domain //IG superfamily
 NT2RP4000925//Fibronectin type III domain
 NT2RP4000927//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP4000973//DnaJ, prokaryotic heat shock protein //Thioredoxins
 45 NT2RP4001079//E1-E2 ATPases
 NT2RP4001080//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP4001117//eubacterial secY protein
 NT2RP4001150//Fibronectin type III domain
 NT2RP4001213//Zinc finger, C2H2 type
 50 NT2RP4001219//Thioredoxins
 NT2RP4001235//Zinc finger, CCHC class
 NT2RP4001433//Zinc finger, C2H2 type
 NT2RP4001498//Ank repeat
 NT2RP4001568//Ank repeat
 55 NT2RP4001644//Eukaryotic protein kinase domain
 NT2RP4001725//WD domain, G-beta repeats
 NT2RP4001753//Zinc finger, C2H2 type
 NT2RP4001790//Zinc finger, C2H2 type

NT2RP2005557//Bacterial mutT protein
 NT2RP2005654//DnaJ, prokaryotic heat shock protein
 NT2RP2005701//Zinc finger, C3HC4 type (RING finger)
 NT2RP2005722//Zinc finger, C2H2 type
 5 NT2RP2005752//TNFR/NGFR cysteine-rich region
 NT2RP2005763//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2005767//HMG (high mobility group) box
 NT2RP2006312//HMG (high mobility group) box
 NT2RP2006464//HMG (high mobility group) box
 10 NT2RP2006571//Cytochrome P450
 NT2RP3000050//Zinc finger, C2H2 type
 NT2RP3000068//PH (pleckstrin homology) domain
 NT2RP3000085//Biotin-requiring enzymes //Carbamoyl-phosphate synthase (CPSase)
 NT2RP3000299//Src homology domain 3
 15 NT2RP3000359//Adenylate kinases
 NT2RP3000366//Ras family (contains ATP/GTP binding P-loop)
 NT2RP3000403//WW/rsp5/WWP domain containing proteins
 NT2RP3000487//WW/rsp5/WWP domain containing proteins
 NT2RP3000512//Homeobox domain
 20 NT2RP3000527//Zinc finger, C2H2 type
 NT2RP3000531//IG superfamily
 NT2RP3000590//Zinc finger, C3HC4 type (RING finger)
 NT2RP3000603//Helix-loop-helix DNA-binding domain
 NT2RP3000605//Zinc finger, C2H2 type
 25 NT2RP3000632//Zinc finger, C2H2 type
 NT2RP3000742//Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 NT2RP3000759//ADP-ribosylation factors (Arf family) (contains ATP/GTP binding P-loop)
 NT2RP3000825//EGF-like domain
 30 NT2RP3000869//ATPases associated with various cellular activities (AAA)
 NT2RP3000994//Double-stranded RNA binding motif
 NT2RP3001057//Zinc finger, C2H2 type
 NT2RP3001084//PH (pleckstrin homology) domain
 NT2RP3001120//Zinc finger, C2H2 type
 35 NT2RP3001140//Thrombospondin type 1 domain
 NT2RP3001150//Forkhead-associated (FHA) domain
 NT2RP3001155//HMG (high mobility group) box
 NT2RP3001214//Zinc finger, C2H2 type
 NT2RP3001268//Zinc finger, C2H2 type
 40 NT2RP3001338//Zinc finger, C2H2 type
 NT2RP3001355//Mitochondrial carrier proteins
 NT2RP3001398//Zinc finger, C2H2 type
 NT2RP3001426//DnaJ, prokaryotic heat shock protein
 NT2RP3001453//ABC transporters
 45 NT2RP3001457//PH (pleckstrin homology) domain
 NT2RP3001472//HMG (high mobility group) box
 NT2RP3001495//Alcohol/other dehydrogenases, short chain type //WW/rsp5/WWP domain containing proteins
 NT2RP3001497//Zinc finger, C3HC4 type (RING finger)
 NT2RP3001724//Helicases conserved C-terminal domain
 50 NT2RP3001792//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP3001943//Zinc finger, C3HC4 type (RING finger)
 NT2RP3001944//Zinc finger, C3HC4 type (RING finger)
 NT2RP3002007//ATPases associated with various cellular activities (AAA)
 NT2RP3002054//Low-density lipoprotein receptor domain class A
 55 NT2RP3002151//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP3002399//MCM2/3/5 family
 NT2RP3002501//Serine/threonine dehydratases
 NT2RP3002602//Thioredoxins

NT2RP2001748//Polyprenyl synthetases
 NT2RP2001756//Zinc finger, C2H2 type
 NT2RP2001839//Eukaryotic protein kinase domain
 NT2RP2001900//Actins
 5 NT2RP2001991//Sodium:neurotransmitter symporter family
 NT2RP2002058//WD domain, G-beta repeats
 NT2RP2002124//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RP2002185//Ubiquitin family
 NT2RP2002208//Zinc finger, C3HC4 type (RING finger)
 10 NT2RP2002256//Cytochrome P450
 NT2RP2002479//ABC transporters
 NT2RP2002503//Zinc finger, C2H2 type
 NT2RP2002520//Ank repeat
 NT2RP2002591//Zinc finger, C2H2 type
 15 NT2RP2002741//Src homology domain 3
 NT2RP2002929//WD domain, G-beta repeats
 NT2RP2002939//Zinc finger, C2H2 type
 NT2RP2002959//Ubiquitin-conjugating enzymes
 NT2RP2002980//Ribosomal protein S10
 20 NT2RP2003137//Ubiquitin family
 NT2RP2003164//Eukaryotic protein kinase domain
 NT2RP2003228//MCM2/3/5 family
 NT2RP2003243//Fibronectin type III domain
 NT2RP2003272//Ubiquitin family
 25 NT2RP2003307//Kinesin light chain repeat
 NT2RP2003401//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases, family 2
 NT2RP2003433//eubacterial secY protein
 NT2RP2003480//Zinc finger, C2H2 type
 NT2RP2003713//Ubiquitin carboxyl-terminal hydrolases family 2
 30 NT2RP2003737//Ubiquitin-conjugating enzymes
 NT2RP2003777//Zinc finger, C3HC4 type (RING finger)
 NT2RP2003840//Ubiquitin-conjugating enzymes
 NT2RP2003857//Ank repeat
 NT2RP2003981//Zinc finger, C3HC4 type (RING finger)
 35 NT2RP2004170//WD domain, G-beta repeats
 NT2RP2004187//Zinc finger, C2H2 type
 NT2RP2004232//Phorbol esters / diacylglycerol binding domain //PH (pleckstrin homology) domain //Eukaryotic
 protein kinase domain
 NT2RP2004389//Ribosomal protein S9
 40 NT2RP2004538//PH (pleckstrin homology) domain
 NT2RP2004568//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2004710//WW/rsp5/WWP domain containing proteins
 NT2RP2004768//Eukaryotic protein kinase domain
 NT2RP2004933//Eukaryotic protein kinase domain
 45 NT2RP2004961//Zinc finger, C2H2 type
 NT2RP2005003//Zinc finger, C3HC4 type (RING finger)
 NT2RP2005012//DnaJ, prokaryotic heat shock protein
 NT2RP2005126//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP2005139//Ank repeat
 50 NT2RP2005140//PH (pleckstrin homology) domain
 NT2RP2005239//Aminotransferases class-V
 NT2RP2005288//Regulator of chromosome condensation (RCC1)
 NT2RP2005293//PH (pleckstrin homology) domain
 NT2RP2005325//Homeobox domain //LIM domain containing proteins
 55 NT2RP2005344//E1-E2 ATPases
 NT2RP2005465//Mitochondrial carrier proteins
 NT2RP2005525//Forkhead-associated (FHA) domain
 NT2RP2005531//Band 4.1 family

NT2RM4002073//AMP-binding enzymes
 NT2RM4002093//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RM4002145//IG superfamily
 NT2RM4002287//Fibronectin type III domain
 5 NT2RM4002527//WD domain, G-beta repeats
 NT2RM4002623//tRNA synthetases class II
 NT2RP1000101//Zinc finger, C2H2 type
 NT2RP1000202//Ank repeat
 NT2RP1000272//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 10 NT2RP1000363//PH (pleckstrin homology) domain
 NT2RP1000376//Ank repeat
 NT2RP1000470//DEAD and DEAH box helicases
 NT2RP1000478//Tubulin
 NT2RP1000522//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 15 NT2RP1000677//Kazal-type serine protease inhibitor domain
 NT2RP1000701//WD domain, G-beta repeats
 NT2RP1000733//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP1000782//4 transmembrane segments integral membrane proteins
 NT2RP1000833//3'-5'-cyclic nucleotide phosphodiesterases
 20 NT2RP1000856//4 transmembrane segments integral membrane proteins
 NT2RP1000947//Ubiquitin-conjugating enzymes
 NT2RP1000959//60s Acidic ribosomal protein
 NT2RP1000966//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP1001033//Tubulin
 25 NT2RP1001080//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RP1001177//Core histones H2A, H2B, H3 and H4
 NT2RP1001247//Transforming growth factor beta like domain
 NT2RP1001294//WD domain, G-beta repeats
 NT2RP1001302//WD domain, G-beta repeats
 30 NT2RP1001313//Heme-binding domain in cytochrome b5 and oxidoreductases
 NT2RP1001457//WD domain, G-beta repeats
 NT2RP1001546//4 transmembrane segments integral membrane proteins
 NT2RP2000008//Zinc finger, C2H2 type
 NT2RP2000040//C2 domain
 35 NT2RP2000045//DnaJ, prokaryotic heat shock protein
 NT2RP2000054//Zinc finger, C3HC4 type (RING finger)
 NT2RP2000070//Cadherin
 NT2RP2000126//Helicases conserved C-terminal domain //SNF2 and others N-terminal domain
 NT2RP2000153//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 40 NT2RP2000224//PH (pleckstrin, homology) domain
 NT2RP2000257//Mitochondrial carrier proteins
 NT2RP2000329//Adenylate kinases
 NT2RP2000414//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP2000448//PH (pleckstrin homology) domain
 45 NT2RP2000660//ATPases associated with various cellular activities (AAA)
 NT2RP2000668//Eukaryotic protein kinase domain
 NT2RP2000710//tRNA synthetases class II
 NT2RP2000764//Aminotransferases class-V
 NT2RP2000842//7 transmembrane receptor (rhodopsin family)
 50 NT2RP2000880//Elongation factor Tu family (contains ATP/GTP binding P-loop)
 NT2RP2000931//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RP2000932//Ank repeat
 NT2RP2001081//C2 domain
 NT2RP2001174//Zinc finger, C2H2 type
 55 NT2RP2001397//Cyclins
 NT2RP2001520//Mitochondrial carrier proteins
 NT2RP2001597//Zinc finger, C3HC4 type (RING finger)
 NT2RP2001740//Ubiquitin carboxyl-terminal hydrolases family 2

NT2RM2000577//tRNA synthetases class I
 NT2RM2000594//C-5 cytosine-specific DNA methylases
 NT2RM2000691//Actins
 NT2RM2000735//Zinc finger, C2H2 type
 5 NT2RM2000740//Helicases conserved C-terminal domain
 NT2RM2000951//FGGY family of carbohydrate kinases
 NT2RM2001324//LIM domain containing proteins
 NT2RM2001499//Amino acid permeases
 NT2RM2001547//DnaJ, prokaryotic heat shock protein //Thioredoxins
 10 NT2RM2001613//eubacterial secY protein
 NT2RM2001670//Zinc finger, C2H2 type
 NT2RM2001700//Acyl-CoA dehydrogenases
 NT2RM2001730//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RM2001813//WD domain, G-beta repeats
 15 NT2RM2001823//Helicases conserved C-terminal domain //SNF2 and others N-terminal domain
 NT2RM2001896//Cytochrome C oxidase subunit II
 NT2RM2001989//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RM2001997//Thioredoxins
 NT2RM2002088//KH domain family of RNA binding proteins
 20 NT2RM2002100//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RM2002109//IG superfamily
 NT2RM4000046//Zinc finger, C3HC4 type (RING finger)
 NT2RM4000104//Zinc finger, C2H2 type
 NT2RM4000167//Kinesin motor domain
 25 NT2RM4000191//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RM4000202//Zinc finger, C2H2 type
 NT2RM4000229//PH (pleckstrin homology) domain
 NT2RM4000344//ATPases associated with various cellular activities (AAA)
 NT2RM4000356//Ras family (contains ATP/GTP binding P-loop)
 30 NT2RM4000471//Aminotransferases class-V
 NT2RM4000496//ATPases associated with various cellular activities (AAA)
 NT2RM4000611//WD domain, G-beta repeats
 NT2RM4000657//C2 domain //Phosphatidylinositol-specific phospholipase C, Y domain
 NT2RM4000712//Ubiquitin carboxyl-terminal hydrolases family 2 //Ubiquitin carboxyl-terminal hydrolases family 2
 35 NT2RM4000733//Forkhead-associated (FHA) domain
 NT2RM4000734//Zinc finger, C2H2 type
 NT2RM4000751//Zinc finger, C2H2 type
 NT2RM4000795//Carboxylesterases
 NT2RM4000996//Zinc finger, C2H2 type
 40 NT2RM4001054//eubacterial secY protein
 NT2RM4001140//Homeobox domain
 NT2RM4001178//DEAD and DEAH box helicases
 NT2RM4001200//Zinc finger, C2H2 type
 NT2RM4001313//Phosphatidylinositol 3- and 4-kinases
 45 NT2RM4001316//Acyl-CoA dehydrogenases
 NT2RM4001320//Src homology domain 3
 NT2RM4001411//PH (pleckstrin homology) domain //Src homology domain 2
 NT2RM4001454//PH (pleckstrin homology) domain
 NT2RM4001483//Zinc finger, C2H2 type
 50 NT2RM4001629//Src homology domain 3
 NT2RM4001758//Eukaryotic protein kinase domain
 NT2RM4001810//Zinc finger, C2H2 type
 NT2RM4001813//Lectin C-type domain short and long forms
 NT2RM4001823//Zinc finger, C2H2 type
 55 NT2RM4001828//Zinc finger, C2H2 type
 NT2RM4001979//Zinc finger, C2H2 type
 NT2RM4001987//IG superfamily
 NT2RM4002013//WD domain, G-beta repeats

HEMBB1001282//Ank repeat
 HEMBB1001294//Ras family (contains ATP/GTP binding P-loop)
 HEMBB1001339//Forkhead-associated (FHA) domain
 HEMBB1001673//Forkhead-associated (FHA) domain //Zinc finger, C3HC4 type (RING finger)
 5 HEMBB1001802//Intermediate filament proteins
 HEMBB1001839//Zinc finger, C2H2 type
 HEMBB1002217//Zinc finger, C2H2 type
 HEMBB1002342//Thioredoxins
 HEMBB1002600//4 transmembrane segments integral membrane proteins
 10 MAMMA1000173//Src homology domain 3
 MAMMA1000388//Zinc finger, C2H2 type
 MAMMA1000402//Reverse transcriptase (RNA-dependent DNA polymerase)
 MAMMA1000612//WD domain, G-beta repeats
 MAMMA1000672//Serine carboxypeptidases
 15 MAMMA1000731//SNF2 and others N-terminal domain
 MAMMA1001008//Eukaryotic aspartyl proteases
 MAMMA1001041//Actinin-type actin-binding domain containing proteins
 MAMMA1001059//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 MAMMA1001105//Zinc finger, C2H2 type
 20 MAMMA1001260//Zinc finger, C3HC4 type (RING finger)
 MAMMA1001576//Tubulin
 MAMMA1001735//Tubulin
 MAMMA1001768//ATPases associated with various cellular activities (AAA)
 MAMMA1001837//Zinc finger, C2H2 type
 25 MAMMA1002170//Ribosomal protein S5
 MAMMA1002385//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 MAMMA1002619//Ubiquitin carboxyl-terminal hydrolases family 2
 MAMMA1002637//Kinesin light chain repeat
 MAMMA1002650//Zinc finger, C2H2 type
 30 MAMMA1002671//AMP-binding enzymes
 MAMMA1002869//LIM domain containing proteins
 MAMMA1002881//SCP-like extracellular Proteins
 MAMMA1002937//Zinc finger, C2H2 type
 MAMMA1002938//Multicopper oxidases
 35 MAMMA1003011//Core histones H2A, H2B, H3 and H4
 MAMMA1003057//WD domain, G-beta repeats
 MAMMA1003127//Myosin head (motor domain) (contains ATP/GTP binding P-loop)
 NT2RM1000086//Zinc finger, C3HC4 type (RING finger)
 NT2RM1000199//CUB domain //Sushi domain
 40 NT2RM1000256//Glutamine amidotransferases class-II
 NT2RM1000499//Ank repeat
 NT2RM1000555//'Cold-shock' DNA-binding domain containing proteins
 NT2RM1000666//'Cold-shock' DNA-binding domain containing proteins //Zinc finger, CCHC class
 NT2RM1000772//WD domain, G-beta repeats
 45 NT2RM1000826//'Cold-shock' DNA-binding domain containing proteins
 NT2RM1000850//Ank repeat //Eukaryotic protein kinase domain
 NT2RM1000852//DEAD and DEAH box helicases //Helicases conserved C-terminal domain
 NT2RM1000882//Heme-binding domain in cytochrome b5 and oxidoreductases
 NT2RM1000885//Zinc finger, C3HC4 type (RING finger)
 50 NT2RM1001059//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 NT2RM1001072//C2 domain //Phosphatidylinositol-specific phospholipase C, X domain //Phosphatidylinositol-specific phospholipase C, Y domain
 NT2RM2000092//Ubiquitin carboxyl-terminal hydrolases family 2
 NT2RM2000101//Zinc finger, C3HC4 type (RING finger)
 55 NT2RM2000191//3'-cyclic nucleotide phosphodiesterases
 NT2RM2000422//Sodium:neurotransmitter symporter family
 NT2RM2000490//C2 domain
 NT2RM2000566//Integrins alpha chain

HEMBA1002177//GATA family of transcription factors //Zinc finger, C2H2 type
 HEMBA1002212//Eukaryotic protein kinase domain
 HEMBA1002215//LIM domain containing proteins
 HEMBA1002419//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 5 HEMBA1002547//Kazal-type serine protease inhibitor domain //Laminin EGF-like (Domains III and V)
 HEMBA1002768//Src homology domain 3
 HEMBA1002810//WW/rsp5/WWP domain containing proteins
 HEMBA1002818//EGF-like domain
 HEMBA1002935//Zinc finger, C2H2 type
 10 HEMBA1002939//Ank repeat
 HEMBA1002973//3'5'-cyclic nucleotide phosphodiesterases
 HEMBA1003077//Fibronectin type III domain
 HEMBA1003250//Eukaryotic protein kinase domain
 HEMBA1003257//Zinc finger, C2H2 type
 15 HEMBA1003281//IG superfamily
 HEMBA1003291//Eukaryotic protein kinase domain
 HEMBA1003433//Forkhead-associated (FHA) domain
 HEMBA1003545//Homeobox domain //LIM domain containing proteins
 HEMBA1003591//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 20 HEMBA1003684//Zinc finger, C2H2 type
 HEMBA1003953//Zinc finger, C2H2 type
 HEMBA1004202//Ras family (contains ATP/GTP binding P-loop)
 HEMBA1004227//Protein phosphatase 2C
 HEMBA1004321//Zinc finger, C2H2 type
 25 HEMBA1004356//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 HEMBA1004408//Peptidyl-prolyl cis-trans isomerases
 HEMBA1004596//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 HEMBA1004734//Ubiquitin-conjugating enzymes
 HEMBA1004973//Fibronectin type III domain
 30 HEMBA1005009//Actins
 HEMBA1005101//RNA recognition motif. (aka RRM, RBD, or RNP domain)
 HEMBA1005581//EGF-like domain //Laminin G domain
 HEMBA1005732//Polyprenyl synthetases
 HEMBA1005737//EF hand
 35 HEMBA1006248//Zinc finger, C2H2 type
 HEMBA1006284//Ubiquitin family
 HEMBA1006293//IG superfamily
 HEMBA1006344//Band 4.1 family
 HEMBA1006445//Ras family (contains ATP/GTP binding P-loop)
 40 HEMBA1006492//Ank repeat
 HEMBA1006559//Zinc finger, C3HC4 type (RING finger)
 HEMBA1006708//WD domain, G-beta repeats
 HEMBA1006737//Ank repeat
 HEMBA1006758//Cadherin
 45 HEMBA1006941//Thioredoxins
 HEMBA1007243//Purine/pyrimidine phosphoribosyl transferases
 HEMBA1007300//3'5'-cyclic nucleotide phosphodiesterases
 HEMBB1000083//IG superfamily
 HEMBB1000317//EGF-like domain //Thrombospondin type 1 domain
 50 HEMBB1000556//Actinin-type actin-binding domain containing proteins //LIM domain containing proteins
 HEMBB1000725//Ras family (contains ATP/GTP binding P-loop)
 HEMBB1000781//Eukaryotic protein kinase domain
 HEMBB1000915//Thrombospondin type 1 domain
 HEMBB1000927//EF hand
 55 HEMBB1000947//Double-stranded RNA binding motif
 HEMBB1001112//eubacterial secY protein
 HEMBB1001175//Ank repeat
 HEMBB1001234//WW/rsp5/WWP domain containing proteins

	PLACE1007243,	PLACE1007274,	PLACE1007282,	PLACE1007317,	PLACE1007375,	PLACE1007386,
	PLACE1007409,	PLACE1007484,	PLACE1007583,	PLACE1007632,	PLACE1007645,	PLACE1007852,
	PLACE1007877,	PLACE1008331,	PLACE1008424,	PLACE1008531,	PLACE1008532,	PLACE1008568,
	PLACE1008715,	PLACE1009045,	PLACE1009319,	PLACE1009338,	PLACE1009368,	PLACE1009493,
5	PLACE1009639,	PLACE1009708,	PLACE1009731,	PLACE1010089,	PLACE1010231,	PLACE1010321,
	PLACE1010622,	PLACE1010811,	PLACE1010917,	PLACE1010954,	PLACE1011090,	PLACE1011214,
	PLACE1011221,	PLACE1011399,	PLACE1011492,	PLACE1011646,	PLACE1011749,	PLACE2000034,
	PLACE2000111,	PLACE2000176,	PLACE2000187,	PLACE2000341,	PLACE2000379,	PLACE2000425,
	PLACE2000458,	PLACE3000020,	PLACE3000218,	PLACE3000226,	PLACE3000244,	PLACE3000413,
10	PLACE4000052,	PLACE4000129,	PLACE4000300,	PLACE4000387,	PLACE4000581,	PLACE4000593,
	PLACE4000650,	THYRO1000394,	THYRO1000395,	THYRO1000570,	THYRO1000748,	THYRO1000756,
	THYRO1001134,	THYRO1001271,	THYRO1001401,	THYRO1001534,	THYRO1001541,	THYRO1001809,
	Y79AA1000258,	Y79AA1000420,	Y79AA1000469,	Y79AA1000734,	Y79AA1000800,	Y79AA1000976,
	Y79AA1001023,	Y79AA1001177,	Y79AA1001394,	Y79AA1001603,	Y79AA1001647,	Y79AA1001846,
15	Y79AA1001874,	Y79AA1002139,	Y79AA1002351,	Y79AA1002399,	Y79AA1002416,	HEMBA1004055,
	HEMBB1001630,	HEMBB1001872,	HEMBB1002044,	HEMBB1002383,	MAMMA1000778,	MAMMA1000859,
	MAMMA1000897,	MAMMA1001073,	MAMMA1002009,	MAMMA1002844,	MAMMA1002947,	MAMMA1003089,
	NT2RM1000092,	NT2RM1000833,	NT2RP2002105,	NT2RP2003668,	NT2RP2006184,	NT2RP3001282,
	NT2RP3002810,	NT2RP3002985,	NT2RP3003059,	NT2RP3003576,	NT2RP3003665,	NT2RP3003799,
20	NT2RP3003828,	NT2RP3003992,	NT2RP3004051,	NT2RP3004155,	OVARC1000890,	OVARC1001117,
	OVARC1001329,	PLACE1001761,	PLACE1002437,	PLACE1004793,	PLACE1005611,	PLACE1005898,
	PLACE1009935,	PLACE1011896,	PLACE2000132,	PLACE2000335,	PLACE3000373,	PLACE3000406,
	PLACE4000250,	PLACE4000487,	PLACE4000494,	THYRO1001320,	THYRO1001537,	THYRO1001828,
	Y79AA1001384					

25 **[0256]** Names of clones whose deduced amino acid sequences were predicted to have functional domains by Pfam search, and names of the matched functional domains are shown below.

When multiple functional domains matched a clone, each domain name was indicated, separated by a double-slash mark, //.

30 HEMBA1000005//DnaJ, prokaryotic heat shock protein
HEMBA1000020//Tubulin
HEMBA1000129//Helicases conserved C-terminal domain
HEMBA1000156//RNA recognition motif. (aka RRM, RBD, or RNP domain)
HEMBA1000158//Fork head domain, eukaryotic transcription factors //Zinc finger, C2H2 type

35 HEMBA1000303//Src homology domain 3 //Zinc finger, C3HC4 type (RING finger)
HEMBA1000411//Ank repeat
HEMBA1000491//Ras family (contains ATP/GTP binding P-loop)
HEMBA1000531//Heat shock hsp70 proteins
HEMBA1000561//Zinc finger, C2H2 type

40 HEMBA1000608//Src homology domain 3
HEMBA1000919//WD domain, G-beta repeats
HEMBA1001043//Ank repeat
HEMBA1001088//LIM domain containing proteins
HEMBA1001137//Zinc finger, C2H2 type

45 HEMBA1001174//ADP-ribosylation factors (Arf family) (contains ATP/GTP binding P-loop)
HEMBA1001247//WW/rsp5/WWP domain containing proteins
HEMBA1001286//Sushi domain
HEMBA1001510//Basic region plus leucine zipper transcription factors
HEMBA1001515//Reverse transcriptase (RNA-dependent DNA polymerase)

50 HEMBA1001661//Cadherin
HEMBA1001723//WD domain, G-beta repeats
HEMBA1001744//Eukaryotic protein kinase domain
HEMBA1001804//Zinc finger, C2H2 type
HEMBA1001819//Zinc finger, C2H2 type

55 HEMBA1001847//Zinc finger, C2H2 type
HEMBA1002035//Bromodomain
HEMBA1002102//Ank repeat
HEMBA1002161//Myosin head (motor domain) (contains ATP/GTP binding P-loop)

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	MAMMA1001030,	MAMMA1001139,	MAMMA1001154,	MAMMA1001388,	MAMMA1001411,	MAMMA1001487,
	MAMMA1001751,	MAMMA1001771,	MAMMA1002461,	MAMMA1002524,	MAMMA1002598,	MAMMA1002684,
	MAMMA1002769,	MAMMA1002890,	MAMMA1002938,	MAMMA1003146,	NT2RM1000035,	NT2RM1000037,
	NT2RM1000062,	NT2RM1000131,	NT2RM1000257,	NT2RM1000260,	NT2RM1000355,	NT2RM1000648,
5	NT2RM1000742,	NT2RM1000800,	NT2RM1000811,	NT2RM1000857,	NT2RM1000867,	NT2RM1000882,
	NT2RM1001008,	NT2RM1001115,	NT2RM1001139,	NT2RM2000259,	NT2RM2000395,	NT2RM2000402,
	NT2RM2000407,	NT2RM2000422,	NT2RM2000566,	NT2RM2000581,	NT2RM2000609,	NT2RM2001370,
	NT2RM2001393,	NT2RM2001499,	NT2RM2001613,	NT2RM2001648,	NT2RM2001659,	NT2RM2001671,
	NT2RM2001718,	NT2RM2001760,	NT2RM2001785,	NT2RM2001823,	NT2RM2001930,	NT2RM2001950,
10	NT2RM2001998,	NT2RM2002049,	NT2RM4000046,	NT2RM4000233,	NT2RM4000433,	NT2RM4000520,
	NT2RM4000634,	NT2RM4000674,	NT2RM4000700,	NT2RM4000764,	NT2RM4000795,	NT2RM4000820,
	NT2RM4000857,	NT2RM4001032,	NT2RM4001054,	NT2RM4001455,	NT2RM4001813,	NT2RM4001930,
	NT2RM4001987,	NT2RM4002054,	NT2RM4002073,	NT2RM4002145,	NT2RM4002146,	NT2RM4002194,
	NT2RM4002339,	NT2RM4002438,	NT2RM4002446,	NT2RM4002452,	NT2RM4002460,	NT2RM4002493,
15	NT2RM4002571,	NT2RP1000191,	NT2RP1000358,	NT2RP1000418,	NT2RP1000547,	NT2RP1000609,
	NT2RP1000677,	NT2RP1000767,	NT2RP1000782,	NT2RP1000856,	NT2RP1001113,	NT2RP1001247,
	NT2RP1001286,	NT2RP1001310,	NT2RP1001311,	NT2RP1001313,	NT2RP1001385,	NT2RP1001449,
	NT2RP1001546,	NT2RP1001569,	NT2RP2000032,	NT2RP2000040,	NT2RP2000070,	NT2RP2000091,
	NT2RP2000114,	NT2RP2000120,	NT2RP2000173,	NT2RP2000175,	NT2RP2000195,	NT2RP2000248,
20	NT2RP2000270,	NT2RP2000283,	NT2RP2000289,	NT2RP2000459,	NT2RP2000516,	NT2RP2000842,
	NT2RP2000892,	NT2RP2001081,	NT2RP2001268,	NT2RP2001295,	NT2RP2001366,	NT2RP2001576,
	NT2RP2001581,	NT2RP2001597,	NT2RP2001947,	NT2RP2001991,	NT2RP2002025,	NT2RP2002312,
	NT2RP2002385,	NT2RP2002479,	NT2RP2002537,	NT2RP2002643,	NT2RP2002701,	NT2RP2002740,
	NT2RP2002857,	NT2RP2003125,	NT2RP2003297,	NT2RP2003433,	NT2RP2003446,	NT2RP2003466,
25	NT2RP2003629,	NT2RP2003777,	NT2RP2003781,	NT2RP2004041,	NT2RP2004194,	NT2RP2004270,
	NT2RP2004681,	NT2RP2004775,	NT2RP2004799,	NT2RP2004936,	NT2RP2005012,	NT2RP2005159,
	NT2RP2005227,	NT2RP2005270,	NT2RP2005344,	NT2RP2005509,	NT2RP2005752,	NT2RP2005781,
	NT2RP2005784,	NT2RP2005812,	NT2RP2006069,	NT2RP2006100,	NT2RP2006141,	NT2RP2006261,
	NT2RP2006571,	NT2RP3000092,	NT2RP3000134,	NT2RP3000333,	NT2RP3000393,	NT2RP3000439,
30	NT2RP3000441,	NT2RP3000531,	NT2RP3000685,	NT2RP3000826,	NT2RP3000852,	NT2RP3001126,
	NT2RP3001176,	NT2RP3001260,	NT2RP3001355,	NT2RP3001383,	NT2RP3001426,	NT2RP3001453,
	NT2RP3001497,	NT2RP3001538,	NT2RP3001716,	NT2RP3001727,	NT2RP3001739,	NT2RP3001799,
	NT2RP3001943,	NT2RP3001944,	NT2RP3002002,	NT2RP3002014,	NT2RP3002054,	NT2RP3002108,
	NT2RP3002163,	NT2RP3002351,	NT2RP3002455,	NT2RP3002549,	NT2RP3002628,	NT2RP3002650,
35	NT2RP3002687,	NT2RP3002701,	NT2RP3002869,	NT2RP3002969,	NT2RP3003008,	NT2RP3003071,
	NT2RP3003101,	NT2RP3003145,	NT2RP3003302,	NT2RP3003353,	NT2RP3003409,	NT2RP3003716,
	NT2RP3003918,	NT2RP3004207,	NT2RP3004454,	NT2RP3004503,	NT2RP4000051,	NT2RP4000151,
	NT2RP4000243,	NT2RP4000259,	NT2RP4000323,	NT2RP4000500,	NT2RP4000560,	NT2RP4000588,
	NT2RP4000713,	NT2RP4000724,	NT2RP4000833,	NT2RP4000878,	NT2RP4000907,	NT2RP4000925,
40	NT2RP4000928,	NT2RP4000973,	NT2RP4000989,	NT2RP4001057,	NT2RP4001064,	NT2RP4001079,
	NT2RP4001117,	NT2RP4001138,	NT2RP4001150,	NT2RP4001174,	NT2RP4001274,	NT2RP4001345,
	NT2RP4001372,	NT2RP4001373,	NT2RP4001379,	NT2RP4001498,	NT2RP4001547,	NT2RP4001571,
	NT2RP4001644,	NT2RP4001677,	NT2RP4001803,	NT2RP4001822,	NT2RP4001975,	NT2RP4002052,
	NT2RP4002075,	NT2RP5003500,	NT2RP5003506,	NT2RP5003522,	NT2RP5003534,	OVARC1000151,
45	OVARC1000241,	OVARC1000335,	OVARC1000700,	OVARC1000722,	OVARC1000751,	OVARC1000850,
	OVARC1000924,	OVARC1000936,	OVARC1000959,	OVARC1000984,	OVARC1001034,	OVARC1001129,
	OVARC1001381,	OVARC1001391,	OVARC1001453,	OVARC1001476,	OVARC1001506,	OVARC1001610,
	OVARC1001702,	OVARC1001703,	OVARC1001713,	OVARC1001745,	OVARC1001767,	OVARC1002127,
	OVARC1002158,	OVARC1002165,	PLACE1000014,	PLACE1000401,	PLACE1000562,	PLACE1000611,
50	PLACE1000656,	PLACE1000712,	PLACE1000909,	PLACE1000948,	PLACE1001241,	PLACE1001257,
	PLACE1001377,	PLACE1001517,	PLACE1001610,	PLACE1001771,	PLACE1001817,	PLACE1001983,
	PLACE1002213,	PLACE1002395,	PLACE1002500,	PLACE1002714,	PLACE1002722,	PLACE1002794,
	PLACE1002851,	PLACE1002908,	PLACE1003045,	PLACE1003238,	PLACE1003296,	PLACE1003369,
	PLACE1003493,	PLACE1003537,	PLACE1003553,	PLACE1003768,	PLACE1003771,	PLACE1003903,
55	PLACE1004197,	PLACE1004258,	PLACE1004270,	PLACE1004289,	PLACE1004473,	PLACE1004743,
	PLACE1004840,	PLACE1004969,	PLACE1005086,	PLACE1005206,	PLACE1005313,	PLACE1005530,
	PLACE1005595,	PLACE1005623,	PLACE1005763,	PLACE1005884,	PLACE1005934,	PLACE1006225,
	PLACE1006754,	PLACE1006901,	PLACE1006935,	PLACE1006956,	PLACE1007014,	PLACE1007111,

EXAMPLE 17

Search for a signal sequence, transmembrane region and functional domain in deduced amino acid sequences

5 **[0252]** The deduced amino acid sequences from the full-length nucleotide sequences were examined to predict the presence of a signal sequence in their amino-termini as well as the presence of a transmembrane region. The amino acid sequences were also searched for a protein functional domain (motif). The examinations for a signal sequence in the amino-terminus, for a transmembrane region and for a functional domain were performed by using PSORT [K. Nakai & M. Kanehisa, Genomics, 14:897-911 (1992)], SOSUI [T. Hirokawa et al., Bioinformatics, 14:378-379 (1998)]
10 (Mitsui Knowledge Industry Co., Ltd.) and Pfam (<http://www.sanger.ac.uk/Software/Pfam/index.shtml>), respectively. When the presence of a signal sequence or a transmembrane region in the amino-terminus was predicted in the amino acid sequence by PSORT or SOSUI, the protein was predicted to be a secretory protein or a membrane protein. When the amino acid sequence matched a functional domain in the Pfam search for a functional domain, the function of the protein is predictable based on the matching data, for example, by referring to the functional categories in PROSITE (<http://www.expasy.ch/cgi-bin/prosite-list.pl>). The functional domain search can be performed by using PROSITE instead of Pfam.

[0253] Search results obtained by using the respective software programs are indicated below.

[0254] Clones whose deduced amino acid sequences were predicted to have signal sequences by PSORT search are as follows:

20 HEMBA1001052, HEMBA1001407, HEMBA1002486, HEMBA1002661, HEMBA1002818, HEMBA1002876,
HEMBA1003086, HEMBA1003711, HEMBA1004752, HEMBA1005991, HEMBA1006067, HEMBA1006173,
HEMBA1006198, HEMBA1006789, HEMBA1006921, HEMBB1000054, HEMBB1000175, HEMBB1002692,
MAMMA1000798, MAMMA1002427, MAMMA1002881, MAMMA1003035, NT2RM1000035, NT2RM1000742,
25 NT2RM1000811, NT2RM1000905, NT2RM1001008, NT2RM2000287, NT2RM2000609, NT2RM2001613,
NT2RM4000634, NT2RM4000778, NT2RM4002339, NT2RM4002460, NT2RP1000782, NT2RP1000856,
NT2RP1001247, NT2RP1001546, NT2RP1001569, NT2RP2001597, NT2RP2002537, NT2RP2004142,
NT2RP2005752, NT2RP2005812, NT2RP3001084, NT2RP3001589, NT2RP3002163, NT2RP3002650,
NT2RP3003145, NT2RP3003242, NT2RP3003621, NT2RP3004282, NT2RP3004503, NT2RP4000051,
NT2RP4000151, NT2RP4000243, NT2RP4000259, NT2RP4000323, NT2RP4000417, NT2RP4001064,
30 NT2RP4001117, NT2RP4001730, NT2RP4001739, NT2RP4002075, NT2RP5003500, OVARC1001154,
PLACE1000611, PLACE1003030, PLACE1003044, PLACE1003369, PLACE1003596, PLACE1004258,
PLACE1005086, PLACE1006239, PLACE1006754, PLACE1006829, PLACE1007954, PLACE1008424,
PLACE1008533, PLACE1008693, PLACE1010622, PLACE1010942, PLACE2000176, PLACE2000341,
PLACE2000379, PLACE2000427, PLACE2000477, PLACE4000431, PLACE4000593, THYRO1000156,
35 THYRO1001134, THYRO1001287, Y79AA1000258, Y79AA1001874, Y79AA1002399, HEMBB1001871,
HEMBB1001925, MAMMA1000778, MAMMA1000897, MAMMA1001080, NT2RP2004300, NT2RP3002985,
NT2RP3003059, OVARC1000689, OVARC1000890, PLACE1005162, PLACE3000399, PLACE3000455,
PLACE4000247, PLACE4000259, PLACE4000494

[0255] Clones whose deduced amino acid sequences were predicted to have transmembrane regions by SOSUI search are as follows:

40 HEMBA1000005, HEMBA1000356, HEMBA1000518, HEMBA1000531, HEMBA1000637, HEMBA1000719,
HEMBA1000817, HEMBA1000822, HEMBA1000870, HEMBA1000991, HEMBA1001052, HEMBA1001085,
HEMBA1001286, HEMBA1001351, HEMBA1001407, HEMBA1001446, HEMBA1001510, HEMBA1001515,
HEMBA1001557, HEMBA1001746, HEMBA1002092, HEMBA1002125, HEMBA1002150, HEMBA1002166,
45 HEMBA1002462, HEMBA1002477, HEMBA1002486, HEMBA1002609, HEMBA1002659, HEMBA1002661,
HEMBA1002780, HEMBA1002818, HEMBA1002876, HEMBA1002921, HEMBA1003077, HEMBA1003079,
HEMBA1003086, HEMBA1003096, HEMBA1003281, HEMBA1003286, HEMBA1003711, HEMBA1003742,
HEMBA1003803, HEMBA1004143, HEMBA1004146, HEMBA1004341, HEMBA1004461, HEMBA1004577,
HEMBA1004637, HEMBA1004752, HEMBA1004756, HEMBA1004850, HEMBA1004889, HEMBA1004923,
50 HEMBA1004930, HEMBA1005029, HEMBA1005035, HEMBA1005050, HEMBA1005552, HEMBA1005588,
HEMBA1005616, HEMBA1005991, HEMBA1006036, HEMBA1006067, HEMBA1006293, HEMBA1006492,
HEMBA1006502, HEMBA1006659, HEMBA1006758, HEMBA1006789, HEMBA1006921, HEMBA1006926,
HEMBA1007203, HEMBB1000050, HEMBB1000054, HEMBB1000556, HEMBB1000593, HEMBB1000631,
HEMBB1000763, HEMBB1000827, HEMBB1000915, HEMBB1000975, HEMBB1001112, HEMBB1001177,
55 HEMBB1001302, HEMBB1001348, HEMBB1001962, HEMBB1002142, HEMBB1002190, HEMBB1002247,
HEMBB1002387, HEMBB1002550, HEMBB1002600, HEMBB1002692, MAMMA1000129, MAMMA1000133,
MAMMA1000277, MAMMA1000278, MAMMA1000410, MAMMA1000416, MAMMA1000472, MAMMA1000714,
MAMMA1000731, MAMMA1000734, MAMMA1000798, MAMMA1000842, MAMMA1000956, MAMMA1001008,

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	HEMBB1000672	F-HEMBB1000672	16122	R-HEMBB1000672	16176
	HEMBB1001197	F-HEMBB1001197	16123	R-HEMBB1001197	16177
	HEMBB1001871	F-HEMBB1001871	16124	R-HEMBB1001871	16178
5	MAMMA1001252	F-MAMMA1001252	16125	R-MAMMA1001252	16179
	MAMMA1002094	F-MAMMA1002094	16126	R-MAMMA1002094	16180
	NT2RM4000634	F-NT2RM4000634	16127	R-NT2RM4000634	16181
	NT2RM4000657	F-NT2RM4000657	16128	R-NT2RM4000657	16182
10	NT2RM4000783	F-NT2RM4000783	16129	R-NT2RM4000783	16183
	NT2RM4000857	F-NT2RM4000857	16130	R-NT2RM4000857	16184
	NT2RM4001178	F-NT2RM4001178	16131	R-NT2RM4001178	16185
	NT2RM4002420	F-NT2RM4002420	16132	R-NT2RM4002420	16186
15	NT2RP2000198	F-NT2RP2000198	16133	R-NT2RP2000198	16187
	NT2RP2000551	F-NT2RP2000551	16134	R-NT2RP2000551	16188
	NT2RP2000660	F-NT2RP2000660	16135	R-NT2RP2000660	16189
	NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
20	NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
	NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
	NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
	NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
25	NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
	NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
	NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
	NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
30	NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
	NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
	NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
	NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
35	NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
	NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
	NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
	NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206
40	NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
	NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
	NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
	NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
45	OVARC1000008	F-OVARC1000008	16157	R-OVARC1000008	16211
	OVARC1000724	F-OVARC1000724	16158	R-OVARC1000724	16212
	OVARC1000751	F-OVARC1000751	16159	R-OVARC1000751	16213
	OVARC1001029	F-OVARC1001029	16160	R-OVARC1001029	16214
50	PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
	PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
	PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
55	PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218

Table 511

5	clone name	name of the 5'-end sequence	SEQ ID NO of the 5'-end sequence	name of the 3'-end sequence	SEQ ID NO of the 3'-end sequence
10	HEMBA1000497	F-HEMBA1000497	16111	R-HEMBA1000497	16165
	HEMBA1001750	F-HEMBA1001750	16112	R-HEMBA1001750	16166
	HEMBA1003854	F-HEMBA1003854	16113	R-HEMBA1003854	16167
15	HEMBA1004193	F-HEMBA1004193	16114	R-HEMBA1004193	16168
	HEMBA1004860	F-HEMBA1004860	16115	R-HEMBA1004860	16169
	HEMBA1005572	F-HEMBA1005572	16116	R-HEMBA1005572	16170
	HEMBA1006038	F-HEMBA1006038	16117	R-HEMBA1006038	16171
20	HEMBA1006092	F-HEMBA1006092	16118	R-HEMBA1006092	16172
	HEMBA1006406	F-HEMBA1006406	16119	R-HEMBA1006406	16173
	HEMBA1006650	F-HEMBA1006650	16120	R-HEMBA1006650	16174
25	HEMBA1006812	F-HEMBA1006812	16121	R-HEMBA1006812	16175

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	NT2RP2002056	F-NT2RP2002056	0. 12
	NT2RP2002677	F-NT2RP2002677	0. 14
5	NT2RP2002755	F-NT2RP2002755	0. 12
	NT2RP2002843	F-NT2RP2002843	0. 11
	NT2RP2003101	F-NT2RP2003101	0. 13
	NT2RP2003799	F-NT2RP2003799	0. 24
10	NT2RP2004095	F-NT2RP2004095	0. 16
	NT2RP2004732	F-NT2RP2004732	0. 18
	NT2RP2004920	F-NT2RP2004920	0. 15
	NT2RP2005454	F-NT2RP2005454	0. 09
15	NT2RP2005776	F-NT2RP2005776	0. 19
	NT2RP2005806	F-NT2RP2005806	0. 27
	NT2RP2005882	F-NT2RP2005882	0. 11
20	NT2RP3001282	F-NT2RP3001282	0. 39
	NT2RP3001723	F-NT2RP3001723	0. 22
	NT2RP3002099	F-NT2RP3002099	0. 20
	NT2RP3003155	F-NT2RP3003155	0. 29
25	NT2RP3004028	F-NT2RP3004028	0. 13
	OVARC1000008	F-OVARC1000008	0. 23
	OVARC1000724	F-OVARC1000724	0. 27
	OVARC1000751	F-OVARC1000751	0. 28
30	OVARC1001029	F-OVARC1001029	0. 25
	PLACE1000814	F-PLACE1000814	0. 21
	PLACE1003030	F-PLACE1003030	0. 26
	PLACE1005549	F-PLACE1005549	0. 16
35	PLACE1007218	F-PLACE1007218	0. 30

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Table 510
The maximal ATGpr1 value of each clone selected in Example 16

clone name	name of sequence	maximal ATGpr1 score
HEMBA1000497	F-HEMBA1000497	0.25
HEMBA1001750	F-HEMBA1001750	0.08
HEMBA1003854	F-HEMBA1003854	0.23
HEMBA1004193	F-HEMBA1004193	0.22
HEMBA1004860	F-HEMBA1004860	0.29
HEMBA1005572	F-HEMBA1005572	0.24
HEMBA1006038	F-HEMBA1006038	0.29
HEMBA1006092	F-HEMBA1006092	0.28
HEMBA1006406	F-HEMBA1006406	0.26
HEMBA1006650	F-HEMBA1006650	0.22
HEMBA1006812	F-HEMBA1006812	0.71
HEMBA1000672	F-HEMBA1000672	0.24
HEMBA1001197	F-HEMBA1001197	0.22
HEMBA1001871	F-HEMBA1001871	0.94
MAMMA1001252	F-MAMMA1001252	0.29
MAMMA1002094	F-MAMMA1002094	0.28
NT2RM4000634	F-NT2RM4000634	0.07
NT2RM4000657	F-NT2RM4000657	0.24
NT2RM4000783	F-NT2RM4000783	0.22
NT2RM4000857	F-NT2RM4000857	0.12
NT2RM4001178	F-NT2RM4001178	0.27
NT2RM4002420	F-NT2RM4002420	0.06
NT2RP2000198	F-NT2RP2000198	0.15
NT2RP2000551	F-NT2RP2000551	0.07
NT2RP2000660	F-NT2RP2000660	0.22
NT2RP2001214	F-NT2RP2001214	0.26
NT2RP2001460	F-NT2RP2001460	0.07
NT2RP2001756	F-NT2RP2001756	0.17

NT2RM4000783, NT2RM4000857, NT2RM4001178, NT2RM4002420, NT2RP2000198, NT2RP2000551, NT2RP2000660, NT2RP2001214, NT2RP2001460, NT2RP2001756, NT2RP2002056, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2003101, NT2RP2003799, NT2RP2004095, NT2RP2004732, NT2RP2004920, NT2RP2005454, NT2RP2005776, NT2RP2005806, NT2RP2005882, NT2RP3001282, NT2RP3001723, NT2RP3002099, NT2RP3003155, NT2RP3004028, OVARC1000008, OVARC1000724, OVARC1000751, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218.

Among them, the following 23 clones was predicted to contain a coding region encoding 100 amino acids or more: HEMBA1000497, HEMBA1003854, HEMBA1004193, HEMBA1006812, HEMBB1001871, NT2RM4000657, NT2RM4001178, NT2RP2001756, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2004095, NT2RP2004920, NT2RP2005806, NT2RP3001282, NT2RP3002099, NT2RP3003155, OVARC1000724, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218. This indicates that the clones encode proteins.

[0247] Table 510 shows maximal ATGpri value determined for each clone. Since the respective maximal ATGpri values for HEMBA1006812, HEMBB1001871 and NT2RRP3001282 are higher than 0.3, the clones would be full-length. Other clones indicated below have maximal ATGpri values of 0.3 or less, and this means that the fullness ratios of the clones are low.

However, the sequences can still be full-length: HEMBA1000497, HEMBA1001750, HEMBA1003854, HEMBA1004193, HEMBA1004860, HEMBA1005572, HEMBA1006038, HEMBA1006092, HEMBA1006406, HEMBA1006650, HEMBB1000672, HEMBB1001197, MAMMA1001252, MAMMA1002094, NT2RM4000634, NT2RM4000657, NT2RM4000783, NT2RM4000857, NT2RM4001178, NT2RM4002420, NT2RP2000198, NT2RP2000551, NT2RP2000660, NT2RP2001214, NT2RP2001460, NT2RP2001756, NT2RP2002056, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2003101, NT2RP2003799, NT2RP2004095, NT2RP2004732, NT2RP2004920, NT2RP2005454, NT2RP2005776, NT2RP2005806, NT2RP2005882, NT2RP3001723, NT2RP3002099, NT2RP3003155, NT2RP3004028, OVARC1000008, OVARC1000724, OVARC1000751, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218

[0248] Table 511 (same as Table 2) shows SEQ ID NOs of the nucleotide sequences located at the 5'-end and 3'-end of each of the 54 clones and the corresponding plasmid clone, which was obtained herein, containing a polynucleotide as an insert. SEQ ID NO for a 5'-end sequence is indicated on the right side of the corresponding Sequence name of 5'-end sequence, and SEQ ID NO for a 3'-end sequence is indicated on the right side of the corresponding Sequence name of 3'-end sequence.

[0249] Swiss-Prot was searched for data homologous to the 5'-end sequences of the selected 54 clones, and GenBank and UniGene were searched for data homologous to the 5'-end and 3'-end sequences of the same clones. The search results are indicated as Homology search results 1-7 in the last part of this SPECIFICATION.

[0250] Based on the matching data obtained by the search, 7 clones presumably encode proteins belonging to any of the categories of secretory or membrane proteins, glycoproteins, signal transduction-associated proteins, transcription-associated proteins, disease-associated proteins, and protein synthesis- and/or protein transport-associated proteins. These were clones exhibiting relatively low homology to any of known proteins belonging to said categories. Here, the term "relatively low homology" means that a nucleotide sequence does not satisfy the conditions under which the nucleotide sequence exhibits "relatively high homology" (which means that, when the nucleotide sequence is compared with the known sequences in Swiss-Prot database, the sequence identity is 60% or higher and the P value is 10^{-10} or less) and that, when the nucleotide sequence is compared with the known sequences in Swiss-Prot database, the sequence to be compared contains 55 nucleotides or more, the sequence identity is 25% or higher, and the P value is 10^{-6} or less.

[0251] Among the 7 clones, clones presumably encoding proteins belonging to the category of secretory or membrane proteins are the two clones, HEMBB1001871 and NT2RM4000857 (which also belong to other categories); clones presumably encoding proteins belonging to the category of glycoproteins are the two clones, HEMBB1001871 and NT2RM4000857 (which also belong to other categories); a clone presumably encoding a protein belonging to the category of signal transduction-associated proteins is PLACE1005549; clones presumably encoding proteins belonging to the category of transcription-associated proteins are the three clones, HEMBA1005572, NT2RP2001756, and NT2RP2005776; a clone presumably encoding a protein belonging to the category of disease-associated proteins is NT2RM4000857 (which also belong to other categories); a clone presumably encoding a protein belonging to the category of protein synthesis- and/or protein transport-associated proteins is HEMBA1001750 (see Examples 12).

Table 509

	Y79AA1002206	7.86	6.79	4.86	2.41	3.6	5.74	5.49	1.51	2.8				
	Y79AA1002208	21.91	17.64	15.14	5.6	4.07	5.57	6.81	5.17	2.55	**	**	-	-
5	Y79AA1002209	14.82	11.28	11.86	6.23	4.7	2.82	4.71	1.33	3.18	**	**	-	-
	Y79AA1002210	13.64	7.39	7.59	9.08	4.62	5.18	20.5	2.05	6.37				
	Y79AA1002211	11.76	19.59	13.47	10.43	6.65	6.52	12.32	8.42	11.25				
	Y79AA1002213	40.78	31.99	22.96	18.41	26.57	14.98	45.88	32.4	41.97				
	Y79AA1002215	54.92	41.69	39.55	24.88	24.36	11.26	37.49	23.6	35.98	*		-	
10	Y79AA1002220	17.03	11.5	20.58	7.13	5.68	5.31	4.57	4.8	6.51	*	*	-	-
	Y79AA1002226	48.55	31.27	31.34	7.35	12.72	13.65	9.19	6.65	11.13	*	**	-	-
	Y79AA1002229	7.88	6.84	5.37	6.02	4.67	2.85	3.67	2.52	3.73	*	*	-	-
	Y79AA1002234	20.83	13.27	12.39	9.34	6.36	3.6	6.9	3.36	5.84	*	*	-	-
	Y79AA1002235	28.03	23.84	21.24	15.07	14.87	9.39	10.75	8.42	13.64	*	**	-	-
15	Y79AA1002246	9.72	14.9	10.35	5.25	6.99	4.12	8.31	3.82	6.69	*		-	
	Y79AA1002258	12.35	12.02	7.88	7.82	11.57	8.55	9.77	4.5	4.81				
	Y79AA1002279	51.52	49.19	41.11	5.28	2.78	2.07	15.99	20.03	22.13	**	**	-	-
	Y79AA1002292	13.64	7.58	4.14	5.73	4.43	2.94	6.45	6.01	8.36				
	Y79AA1002298	9.43	8.29	4.77	3.29	5.48	4.2	4.42	4.85	3.79				
20	Y79AA1002307	9.31	8.07	6.62	3.9	4.4	2.44	5.17	1.76	2.58	*	*	-	-
	Y79AA1002309	8.88	7.96	8.79	3.32	4.26	2.83	4.23	2.88	3.97	**	**	-	-
	Y79AA1002311	16.51	9.27	11.76	6.9	6.2	3.45	7.45	3.09	6.4	*		-	-
	Y79AA1002334	13.05	8.36	7.7	5.13	4.89	3.36	5.91	4.32	5.92	*		-	-
	Y79AA1002351	13.61	12.49	9.42	7.1	4.15	6.19	7.95	4.93	5.8	*	*	-	-
	Y79AA1002355	31.74	30.6	21.85	12.21	15.81	9.74	20.54	18.29	18.48	*	*	-	-
25	Y79AA1002361	23.42	15.4	18.02	12.53	10.73	6.85	25.86	17.1	25.5	*		-	
	Y79AA1002365	12.42	6.37	7.19	3.15	4.11	3.03	4.29	4.74	4.01				
	Y79AA1002373	8.95	6.89	5.46	5.13	4.81	3.4	9.6	4.57	7.84				
	Y79AA1002376	1550.5	2569.2	1680.8	462.62	827.86	616.71	1477.6	1040.3	1062.1	*		-	-
	Y79AA1002378	20.24	17.32	13.54	5.14	9.41	4.23	19.28	11.07	16.88	*		-	-
30	Y79AA1002381	116.11	128.86	74.48	110.66	141.78	92.68	155.95	123.08	170.94				
	Y79AA1002388	33.4	33.3	27.31	13.85	26.75	11.62	21.29	16.32	21.24		*	-	-
	Y79AA1002399	11.13	8.22	7.72	4.28	5.54	4.87	7.56	5.9	6.25	*		-	-
	Y79AA1002407	12.66	14.43	18.13	7.72	14.18	6.84	5.83	9.78	4.59	*	*	-	-
	Y79AA1002413	16.98	12.77	14.95	6.14	9.13	4.62	8.44	10.73	7.99	*	*	-	-
	Y79AA1002416	7.52	8.19	8.76	5.47	10.72	5.8	8.2	6.05	6.59				
35	Y79AA1002429	17.73	18.61	8.81	5.82	10.24	4.73	3.65	6.89	5.66	*		-	-
	Y79AA1002431	3.38	3.05	6.2	3.01	5.89	1.6	2.81	2.79	1.69				
	Y79AA1002433	9.94	11.67	9.29	5.11	5.57	3.18	3.49	4.6	3.87	**	**	-	-
	Y79AA1002445	33.47	25.62	23.49	15.99	10.67	7.02	18.92	25.26	13.87	*		-	-
	Y79AA1002461	7.94	6.22	7.84	3.36	7.35	4.7	3.49	2.25	3.85		**	-	-
	Y79AA1002466	778.44	339.4	681.02	542.56	499.15	369	592.67	971	768.71				
40	Y79AA1002471	11.38	8.13	15.35	12.81	13.4	11.43	4.94	6.06	4.47	*		-	-
	Y79AA1002472	31.22	33.06	31.17	18.15	21.85	9.34	16.29	20.14	20.03	*	**	-	-
	Y79AA1002474	10.68	12.29	10.71	6.77	7.3	7.75	3.17	7.37	4.86	**	*	-	-
	Y79AA1002482	30.09	33.68	36.63	19.02	23.45	17.38	21.9	25.81	23.08	**	*	-	-
	Y79AA1002487	8.33	8.29	7.43	7.28	8.45	6.44	5.34	3.78	3.86		**	-	-
45	Y79AA1002490	143.18	106.89	117.63	56.22	71.49	57.31	59.76	51.39	52.37	**	**	-	-
	Y79AA1002493	44.75	41.56	40.36	20.64	28.52	19.33	38.02	46.19	46.7	**		-	-
	ZRV6C1006278	5.26	7	5.52	3.16	2.97	2.19	2.99	2.6	2.72	**	**	-	-

EXAMPLE 16

Selection of novel cDNA clones from cDNA libraries prepared by oligo-capping method

[0246] The following 54 clones were newly selected from cDNA libraries prepared by oligo-capping method, based on the criterion that the 5'-end sequence of a cDNA clone contained a coding region which was initiated with ATG codon and which encoded 50 amino acids or more:

HEMBA1000497, HEMBA1001750, HEMBA1003854, HEMBA1004193, HEMBA1004860, HEMBA1005572, HEMBA1006038, HEMBA1006092, HEMBA1006406, HEMBA1006650, HEMBA1006812, HEMBB1000672, HEMBB1001197, HEMBB1001871, MAMMA1001252, MAMMA1002094, NT2RM4000634, NT2RM4000657,

Table 508

	Y79AA1001787	5.04	3.62	6.16	2.83	4.49	3.23	3.33	2.97	2.01			
	Y79AA1001793	120.15	97.84	90.95	70.69	108.02	70.38	56.01	50.79	40.18	**	*	-
	Y79AA1001795	5.66	5.4	6.21	4.2	5.9	3.1	3.91	2.49	2.28	**	*	-
5	Y79AA1001799	15.09	14.95	16.22	8.14	9.02	8.13	5.96	8.67	5.96	**	**	-
	Y79AA1001800	25.64	16.41	26.8	14.68	31.06	13.67	17.04	13.04	14.7	*	*	-
	Y79AA1001801	18.54	13.81	14.73	8.69	14.72	7.61	10.02	6.35	8.81	*	*	-
	Y79AA1001803	4.48	5.85	6.44	4.5	2.73	1.98	4.84	2.96	2.69			-
	Y79AA1001805	40.28	26.89	34.51	17.56	11.83	10.76	12.11	14.87	11.59	**	**	-
10	Y79AA1001807	42.38	40.49	41.45	39.87	44.56	37.42	30.19	22.76	22.12	**	**	-
	Y79AA1001827	9.32	11.28	9.81	8.01	8.55	6.68	5.41	8.86	5.83	*	*	-
	Y79AA1001846	22.14	19.33	18.59	11.82	17.37	13.61	12.56	15.77	9.7	*	*	-
	Y79AA1001848	8.04	9.99	9.11	5.33	6.98	5.01	6.65	6.63	5.92	*	*	-
	Y79AA1001853	10.87	10.1	9.06	5.77	8.29	4.56	4.63	5.75	4.91	*	**	-
	Y79AA1001863	24.97	27.27	22.17	17.04	22	11.97	13.95	21.02	18.07	*	*	-
15	Y79AA1001866	13.46	10.72	11.93	6.28	4.52	4.93	6.56	4.65	4.51	**	**	-
	Y79AA1001874	2.1	1.41	2.93	0.78	1.08	0.95	1.46	1.22	1.08			-
	Y79AA1001875	31.78	22.74	37.59	12.11	19.59	16.03	9.16	15	14.61	*	*	-
	Y79AA1001907	517.52	778.32	567.93	750.86	875.67	821.68	479.34	605.65	529.63			-
	Y79AA1001908	5.27	6.69	9.96	3.58	5.35	3.25	2.93	3.09	1.48	*	*	-
	Y79AA1001923	8.48	16.75	14.8	9.28	7.65	6.34	6.48	6.75	5.36	*	*	-
20	Y79AA1001927	18.05	21.6	19.1	12.29	16.25	12.12	14.45	14.01	14.79	*	**	-
	Y79AA1001930	12.23	24.48	15.86	10.91	18.16	6.72	4.54	5.11	5.37	*	*	-
	Y79AA1001932	61.74	56.58	52.19	37.44	49.06	33.54	58.52	51.29	51.59	*	*	-
	Y79AA1001933	9.52	6.51	10.88	6.04	6.05	6.44	4.27	11.51	4.91			-
	Y79AA1001942	10.76	6.66	7.93	7.97	5.94	4.4	3.29	2.7	3.99	*	*	-
	Y79AA1001963	138.12	106.9	117.87	95.52	130.54	91.9	83.12	67.08	69.04	*	*	-
25	Y79AA1001968	147.27	91.21	62.32	82.32	49.63	76.26	20.14	2.41	10.4	*	*	-
	Y79AA1001983	12.1	16.25	14.99	5.06	8.59	7.83	3.08	3.73	3.02	*	**	-
	Y79AA1002000	8.87	9.86	16.46	6.28	9.55	3.45	4.66	4.84	4.51	*	*	-
	Y79AA1002004	46.17	58.66	37.42	34.61	47	15.44	17.13	16.13	8.67	**	*	-
	Y79AA1002008	16.52	21.1	20.83	11.85	16.14	12.02	7.95	15.74	18.89	*	*	-
	Y79AA1002012	13.85	12.54	10.41	5.85	6.84	6.65	8.76	11.21	9.52	**	*	-
30	Y79AA1002017	15.21	10.94	15.42	6.73	12.23	8.22	1.29	15.64	2.06			-
	Y79AA1002022	42.57	32.77	37.11	21.13	22.85	16.91	21.09	21.59	19.22	**	**	-
	Y79AA1002027	10.35	8.04	15.25	2.11	4.6	4.84	5.16	2.67	4.2	*	*	-
	Y79AA1002050	13.21	11.11	14.39	7.26	8.9	5.9	6.9	7.16	7.33	*	**	-
	Y79AA1002058	167.29	130.21	156.91	193.04	186.39	149.35	127.37	93.56	137.15			-
	Y79AA1002080	54.35	76.53	61.68	36.08	41.21	29.16	24.43	9.99	5.28	*	**	-
35	Y79AA1002082	49.46	32.53	50.05	16.65	20.72	17.6	35.11	28.1	35.48	*	*	-
	Y79AA1002085	113.45	69.34	64.62	65.01	102.96	72.96	82.29	90.04	44.91			-
	Y79AA1002087	33.46	37.19	43.9	21.85	20.1	22.68	15.4	10.55	10.37	**	**	-
	Y79AA1002089	5.33	3.94	7.12	3.04	1.49	2.47	3.1	3.24	2.29	*	*	-
	Y79AA1002070	67.39	142.78	83.33	77.38	149.87	86.41	153.3	88.92	78.17			-
	Y79AA1002074	1225	1102	498.16	721.59	1162.7	771.32	764.45	491.49	312.45			-
40	Y79AA1002076	8.24	11.39	15.58	4.01	6.13	3.99	13	9.01	11.16	*	*	-
	Y79AA1002083	11.94	6.94	9.48	4.49	4.53	3.15	4.48	1.37	1.78	*	*	-
	Y79AA1002084	20.05	13.84	17.88	7.41	8.36	6.24	10.42	8.8	8.13	**	*	-
	Y79AA1002086	15.57	8.01	9.58	4.16	3.65	3.41	3.6	4.3	3.59	*	*	-
	Y79AA1002087	265.36	258.38	345.6	296.44	355.85	246.9	347.62	322.78	317.34			-
	Y79AA1002089	15.8	12.04	15.08	6.67	9.17	6.14	12.29	9.32	9.91	**	*	-
45	Y79AA1002093	13.66	12.64	9.33	5.35	6.21	4.47	3.01	2.8	2.57	**	**	-
	Y79AA1002101	6.58	7.04	8.08	3.07	2.77	1.96	3.3	4.71	1.2	**	*	-
	Y79AA1002103	11.98	11.79	10.9	4.11	5.96	4.92	4.54	2.99	2.4	**	**	-
	Y79AA1002115	15.51	18.46	14.58	9.3	11.82	8.45	4.13	9.04	5.86	*	**	-
	Y79AA1002121	6.93	5.28	6.92	6.7	3.27	4.36	3.68	3.01	3.29	**	*	-
50	Y79AA1002125	40.85	21.04	26.11	14.22	18.4	17.66	12.21	8.56	11.75	*	*	-
	Y79AA1002129	7.76	7.25	11.87	4.36	3.94	3.2	3.76	3.8	4.37	*	*	-
	Y79AA1002131	5.89	3.57	4.65	1.58	2.79	2.27	3.32	1.61	2.75	*	*	-
	Y79AA1002139	8.48	7.22	6.23	4.06	4.42	2.47	3.44	3.86	2.67	*	**	-
	Y79AA1002144	53.23	47.33	47.13	32.96	42.65	28.86	30.69	36.28	25.23	*	**	-
	Y79AA1002177	14.09	13.17	13.65	5.87	6.08	5.05	5.82	4.1	4.34	**	**	-
55	Y79AA1002183	76.21	99.47	69.64	24.87	40.12	25.67	26.7	15.32	11.38	**	**	-
	Y79AA1002202	29.69	18.4	18.04	8.22	10.65	9.82	13.41	13.22	16.71	*	*	-
	Y79AA1002204	4.17	2.23	3.23	5.04	1.05	2.68	3.28	2.4	2.69			-

Table S07

	Y79AA1001211	21.15	21.66	22.83	13.08	12.91	10.43	0.44	20.18	19.17	**	-	-
	Y79AA1001212	128.68	116.33	107.84	151.9	151.84	105.91	80.87	83.57	67.14	**	-	-
5	Y79AA1001216	31.68	27.07	33.84	18.3	19.09	14.63	17.05	18.24	23.12	**	*	-
	Y79AA1001228	7.97	5.25	12.95	3.35	3.4	3.64	3.41	10.17	4.89			
	Y79AA1001233	8.26	6.53	16.79	3.45	3.77	2.51	2.28	3.79	3.13			
	Y79AA1001236	8.84	5.25	11.14	5.02	5.53	3.13	4.08	4.7	2.24			
	Y79AA1001239	74.22	59.48	53.54	45.62	83.11	43.05	50.35	42.44	46.18			
	Y79AA1001240	70.83	49.07	64.4	30.83	35.04	20.32	48.49	59.48	54.67	*	-	-
10	Y79AA1001255	8.89	10.29	10.55	7.47	6.78	6.42	0.72	6.35	0.76	**	*	-
	Y79AA1001264	14.2	10.19	13.36	9.71	10.05	8.67	4.58	4.84	6.39	**	-	-
	Y79AA1001272	15.16	12.6	13.5	3.77	6.76	5.59	8.86	7.59	6.53	**	**	-
	Y79AA1001281	8.87	7.72	6.68	4.17	4.8	3.47	2.65	2.22	2.28	**	**	-
	Y79AA1001299	9.39	10.08	9.6	4.21	5.3	4.49	3.28	3.68	3.29	**	**	-
	Y79AA1001312	40.47	42.72	35.81	17.92	27.92	14.77	29.7	20.85	26.67	*	*	-
15	Y79AA1001319	31.92	23	20.23	23.19	28.4	16.96	30.43	20.02	18			
	Y79AA1001323	32.99	27.82	34.32	14.07	19.03	14.31	25.9	23.32	27.9	**	-	-
	Y79AA1001328	28.33	16.1	28.16	14.56	12.93	12.69	4.94	7.51	4.42	*	-	-
	Y79AA1001343	12.56	8.01	10.99	6.43	7.59	7.49	6.83	9.08	8.27			
	Y79AA1001351	6.64	4.21	8.23	2.99	1.8	1.8	2.08	2.52	1.68	*	*	-
20	Y79AA1001364	8.06	5.02	6.89	3.26	2.98	2.12	4.06	2.93	1.74	*	*	-
	Y79AA1001367	145.27	118.42	143.46	117.7	132.8	94.77	76.65	57.61	66.99	**	-	-
	Y79AA1001384	45.86	52.17	41.89	27.43	34.83	21.42	31.33	29.86	26.76	*	**	-
	Y79AA1001391	10.96	12.22	10.34	2.99	9.55	4.77	7.43	7.31	6.01	**	-	-
	Y79AA1001394	24.73	14.05	14.28	9.7	6.32	9.07	6.42	10.16	9.01			
	Y79AA1001402	12.42	7.77	9.12	5.62	4.8	4.67	4.74	3.87	4.52	*	*	-
	Y79AA1001410	5.24	2.77	3.09	3.28	1.53	1.95	2.62	0.78	1.99			
25	Y79AA1001414	27.49	18.53	20.44	10.78	9.47	10.2	16.16	13.43	11.92	*	*	-
	Y79AA1001426	271.77	221.23	225.78	200.19	221.07	217.77	202.63	211.55	197.64			
	Y79AA1001427	15.18	10.97	10.9	6.77	10.81	5.39	6.71	6.07	6.74	*	-	-
	Y79AA1001430	22.25	17.58	19.52	10.25	13.63	7.85	11.32	11.76	10.84	*	**	-
	Y79AA1001439	6.75	4.58	4.31	2.6	4.23	1.25	2.94	2.35	2.98	*	-	-
	Y79AA1001485	17.57	9.99	14.92	13.26	10.56	8.25	5.32	10.37	14.36			
30	Y79AA1001493	20.47	12.23	13.25	9.02	11.27	6.64	13.37	13.83	16.06			
	Y79AA1001511	14.52	8.05	9.32	8.33	8.58	5.82	5.2	6	5.96			
	Y79AA1001523	10.03	6.72	7.33	6.48	4.81	3.84	4.54	3.21	2.59	*	-	-
	Y79AA1001530	11.44	7.57	8.59	6.64	8.84	5.29	6.2	7.76	5.77			
	Y79AA1001532	13.55	14.73	11.04	5.35	5.16	4.93	5.08	4.52	6.49	**	**	-
35	Y79AA1001533	56.5	68.51	50.9	29.97	36.82	26.92	69.28	61.19	56.58	*	-	-
	Y79AA1001541	8.12	8.52	5.25	4.02	4.89	2.24	4.69	2.24	2.18	*	*	-
	Y79AA1001548	67.38	47.55	44.48	30.61	26.42	21.92	46.07	51.22	51.86	*	-	-
	Y79AA1001555	14.35	9.75	8.13	8.86	4.86	4.97	8.19	7.68	10.98			
	Y79AA1001562	18.44	13.73	12.82	8.99	10.6	7.19	10.2	9.39	15.75	*	-	-
	Y79AA1001581	24.34	18.91	17.62	9.8	12.04	10.4	15.9	9.86	16.33	*	-	-
40	Y79AA1001585	30.32	20.83	22.83	20.22	25.12	15.45	0.69	15.13	0.14	*	-	-
	Y79AA1001592	28.53	25.65	26.36	29.33	16.66	7.28	20.69	8.44	14.72	*	-	-
	Y79AA1001594	23.28	21.85	24.28	14.25	15.59	10.11	22.55	14.53	20.27	**	-	-
	Y79AA1001603	12.47	11.04	10.84	6.8	11.48	6.78	22.63	14.33	27.5			
	Y79AA1001613	25.09	18.27	18.81	23.47	16.84	13.34	20.3	16.35	18.99			
	Y79AA1001630	22.94	13.91	12.22	13.14	13.53	8.02	11.73	12.98	17.67			
45	Y79AA1001647	33.54	24.86	22.66	23.22	25.84	16.49	26.93	21.3	26.55			
	Y79AA1001664	12.2	10.56	13.13	7.78	6.05	5.54	10.34	8.1	11.69	**	-	-
	Y79AA1001665	11.52	9	9.75	6.87	7.31	6.13	5.92	6.12	7.69	*	*	-
	Y79AA1001679	24.51	18.98	17.88	11.13	12.84	9.27	24.02	13.28	18.8	*	-	-
	Y79AA1001692	7.23	6.72	8.1	2.24	2.83	2.63	3.64	2.31	3.06	**	**	-
	Y79AA1001696	14.23	12.31	9.21	5.29	8.48	4.61	14.27	13.28	13.01	*	-	-
50	Y79AA1001705	12.04	11.4	8.55	9.76	11.8	6.21	4.61	5.98	4.57	**	-	-
	Y79AA1001711	13.72	7.72	10.63	6.49	4.11	3.08	6.08	4.06	7.4	*	-	-
	Y79AA1001717	4.71	1.55	3.26	3.34	1.71	0.7	1.49	1.63	1.39			
	Y79AA1001719	22.12	12.72	16.33	9.89	12.72	6.53	12.21	11.1	11.76			
	Y79AA1001727	38.96	27.16	22.35	13.65	17.35	9.31	38.01	28.39	34.23	*	-	-
	Y79AA1001750	21.5	17.52	18.42	10.15	15.4	8.01	11.94	9.9	11.41	*	**	-
55	Y79AA1001760	127.24	115.03	77.78	58.28	73.6	43.62	140.25	100.84	138.73	*	-	-
	Y79AA1001777	6.77	6.32	6.37	2.57	5.24	2.46	5.26	4.15	4.71	*	**	-
	Y79AA1001781	2.86	2.24	2.15	1.61	1.86	2.55	0.74	1.53	1.3	*	-	-

Table 506

	Y79AA1000778	20.06	13.62	11.66	10.8	9.69	8.29	13.52	13.51	12.96			
	Y79AA1000782	11.43	7.55	9.19	7.76	8.46	7.38	6.62	5.21	4.82	*		
5	Y79AA1000784	13.14	10.21	12.84	4.67	6.08	6.71	10.7	9.6	9.71	**		
	Y79AA1000794	46.19	39.88	38.82	22.35	24.71	16.75	37.28	33.24	34.17	**		
	Y79AA1000800	7.18	3.91	5.09	0.48	0.36	0.17	2.34	0	0	**	*	
	Y79AA1000802	11.45	13.12	10.85	5.78	8.8	4.96	3.89	4.75	3.77	*	**	
	Y79AA1000805	21.22	18.33	12.77	6.7	11.14	4.1	15.87	12.52	16.72	*		
10	Y79AA1000814	27.49	21.74	19.65	16.59	17.35	12.96	22	17.22	22.17			
	Y79AA1000823	14.24	7.73	9.85	4.94	4.36	5.06	6.73	6.96	8.36	*		
	Y79AA1000824	16.24	10	9.76	8.11	9.28	6.16	6.82	5.83	8.72			
	Y79AA1000827	9.71	3.82	4.6	3.63	3.22	2.56	3.51	2.06	2.83			
	Y79AA1000831	7.38	5.79	7.27	4.13	3.32	2.68	4.43	0.57	1.26	**	*	
	Y79AA1000833	22.35	19.25	16.88	12.38	7.9	8.89	19.38	11.86	18.04	**		
	Y79AA1000850	33.44	40.57	26.99	19.52	21.81	14.62	28.93	23.5	32.01	*		
15	Y79AA1000856	6.26	7.4	4.9	2.81	5.29	2.12	4.31	2.48	3.63	*		
	Y79AA1000862	8.84	4.04	5.05	7.04	4.31	3.07	5.61	5.34	5.05			
	Y79AA1000876	21.17	10.25	11.73	7.67	6.82	6.01	5.79	7.01	6			
	Y79AA1000888	32.02	25.94	21.68	17.64	17.29	12.59	18.41	20.33	23.17	*		
	Y79AA1000902	6.26	5.68	6.17	4.27	4.4	2.72	2.73	2.97	2.19	*	**	
20	Y79AA1000935	76.11	52.08	52.32	70.65	71.58	69.22	39.77	35.2	32.3	*		
	Y79AA1000958	48.22	40.7	37.08	25.45	33.48	19.11	26.33	21.44	21.93	*	**	
	Y79AA1000962	46.82	45.31	38.99	22.79	24.29	17.48	60.82	38.12	53.14	**		
	Y79AA1000963	25.33	35.17	19.37	13.71	17.32	12.79	14.75	14.39	13.92			
	Y79AA1000966	41.31	37.86	31.25	27.6	23.69	23.7	32	18.68	25.22	*		
	Y79AA1000967	19.13	9.69	10.03	8.88	7.99	4.65	1.27	8.2	0.25			
25	Y79AA1000968	25.81	16.48	14.41	11.57	12.79	6.94	25.2	19.6	28.68			
	Y79AA1000969	40.02	39.97	33.37	23.55	39.74	36.04	32.93	30.52	18.66			
	Y79AA1000976	22.75	15.17	16.99	14.98	19.84	15.88	14.85	10.98	11.26			
	Y79AA1000978	8.83	7.09	6.8	4.57	5.39	2.66	4.35	3.43	4.33	*	**	
	Y79AA1000985	8.02	3.9	5.83	4.32	4.39	1.76	4.37	2.17	1.99			
	Y79AA1000989	295.68	243.92	218.77	78.52	138.6	77.87	70.03	46.49	61.43	**	**	
30	Y79AA1000991	12.26	8.35	10.64	5.63	5.93	8.76	7.23	8.65	6.01			
	Y79AA1001013	42.37	37.07	51.31	24.72	27.17	18.23	24.27	34.59	22.31	*	*	
	Y79AA1001014	16.53	17.99	19.41	12.93	20.24	14.74	13.35	18.92	15.51			
	Y79AA1001019	8.07	6.21	8.29	4.14	6.05	5.04	2.84	5.3	3.51	*	*	
	Y79AA1001020	13.89	16.72	13.86	21.03	12.53	10.76	9.5	10.16	11.46	*		
	Y79AA1001023	7.28	6.24	9.28	4.31	8.03	5.29	3.01	3.87	2.51	*		
35	Y79AA1001030	28.41	43.95	33.86	36.62	21.08	22.74	16.86	22.85	18.21	*		
	Y79AA1001035	10.6	13.73	10.93	9.68	10.73	7.69	7.26	7.72	8.91	*		
	Y79AA1001041	14.5	11.14	15.3	8.73	9.98	7.28	9.17	13.51	15.31	*		
	Y79AA1001043	22.79	19.12	22.36	8.78	13.48	10.29	9.42	13.86	17.78	**	*	
	Y79AA1001048	12.86	13.5	12.88	6.65	11.19	7.59	6.44	7.55	6.3	*	**	
	Y79AA1001056	6.67	9.77	12.37	5.42	9.5	6.82	2.1	3.99	2.72	*		
40	Y79AA1001061	11.12	17.1	15.18	5.75	10.72	7.78	8.04	10.36	12.77	*		
	Y79AA1001062	4.7	4.96	6.24	2.95	3.71	2.56	2.92	2.19	2.28	*	**	
	Y79AA1001068	6.22	3.59	5.75	4.34	5.98	4.49	3.29	3.38	3.07			
	Y79AA1001073	17.6	17.56	20.08	5.98	13.24	7.19	16.17	13.71	20.42	*		
	Y79AA1001077	7.3	2.86	6.26	7.08	4.43	3.45	1.86	5.22	1.86			
	Y79AA1001078	7.29	5.05	6.25	3.73	4.63	2.99	3.3	5.06	3.66	*		
45	Y79AA1001081	7.84	5.5	8.85	3.05	3.81	3.52	2.8	3.1	3.74	*	*	
	Y79AA1001088	36.17	40.23	42.34	24.5	22.27	19	24.36	30.94	32.13	**	*	
	Y79AA1001089	332.88	366.62	308.34	284.36	390.54	406.65	237.36	376.46	392			
	Y79AA1001090	7.05	4.26	8.66	4.12	3.36	2.71	3.39	1.8	2.5	*		
	Y79AA1001095	22.53	22.64	23.57	8.51	14.07	9.94	17.11	15.32	15.62	**	**	
50	Y79AA1001142	114.5	98.72	131.88	102.55	114.58	143.24	69.56	60.81	74.51	*		
	Y79AA1001145	98.33	98.57	100.63	125.36	121.38	91.71	78	59.2	74.76	**		
	Y79AA1001162	126.48	83.55	107.65	92.38	93.62	73.32	97.56	124.59	105.84			
	Y79AA1001167	12.61	7.85	13.06	7.92	8.62	6.56	7.15	16.96	9.58			
	Y79AA1001176	75.12	80.1	84.48	76.52	65.54	64.34	46.71	59.03	47.45	**		
	Y79AA1001177	8.46	7.48	10.23	4.06	5.51	3.51	3.45	4.68	2.96	*	**	
	Y79AA1001179	54.1	53.61	67.63	28.64	33.09	26.42	13.82	47.33	10.91	**		
55	Y79AA1001185	32.79	30.94	36.04	15.68	21.34	16.32	19.9	17.65	25.06	**	**	
	Y79AA1001201	45.08	50.33	41.06	13.71	31.35	17.81	48.16	43.78	38.95	*		
	Y79AA1001205	18.73	12.88	21.83	8.38	8.46	6.14	7.31	11.86	11.81	*		

Table 505

	Y79AA1000285	22.27	16.31	18.63	6.85	10.3	7.36	15.47	22.31	18.9	**	-	-
	Y79AA1000295	4.35	2.68	5.01	1.59	3.76	2.11	2.3	2.59	2.22		-	-
5	Y79AA1000307	6.82	4.93	8.12	6.39	9.3	7.36	3	5.88	4.49		-	-
	Y79AA1000313	4.28	2.34	3.94	1.66	3.1	3.25	1.93	2.47	1.62		-	-
	Y79AA1000314	1.57	1.83	4.11	1.4	1.41	0.98	0.87	0.14	1.73		-	-
	Y79AA1000328	3.78	3.39	5.43	2.09	10.7	2.68	3.27	3.33	3.32		-	-
	Y79AA1000334	9.59	9.06	7.49	5.79	6.64	3.5	5.12	3.62	4.02	* **	-	-
	Y79AA1000342	17.48	13.29	18.31	13.02	15.9	13.67	34.2	31.79	30.43	**	-	+
10	Y79AA1000346	17.08	11.48	18.69	8.72	9.82	6.74	5.35	5.97	4.15	* **	-	-
	Y79AA1000347	28.82	26.71	31.26	22.14	31.96	28.74	20.9	20.82	19.69	**	-	-
	Y79AA1000349	15.95	20.35	20.7	17.44	19.4	18.58	5.37	7.08	5.15	**	-	-
	Y79AA1000355	11.9	11.6	17.89	14.98	13.32	9.11	5.25	6.54	7.95	*	-	-
	Y79AA1000368	5.31	4.56	7.88	4.89	6.47	3.77	2.77	3.21	1.95	*	-	-
	Y79AA1000388	9.85	11.63	11.71	9.62	9.75	4.51	5.03	4.57	5.04	**	-	-
15	Y79AA1000392	13.05	13.68	11.25	7.92	12.06	6.34	12.1	13.67	13.52		-	-
	Y79AA1000405	100.68	62.31	79.28	103.67	109.59	81.35	71.19	52.24	66.44		-	-
	Y79AA1000410	8.13	6.1	10.22	6.13	5.36	5.28	4.83	4.8	5.17		-	-
	Y79AA1000420	11.1	11.77	13.46	5.41	8.52	5.43	6.77	7.21	5.64	* **	-	-
	Y79AA1000423	7.61	9.21	8.96	4.46	6.29	5.4	4.36	2.75	3.4	* **	-	-
20	Y79AA1000426	5.75	6.96	10.05	4.68	5.08	3.97	3.38	2.69	2.24	*	-	-
	Y79AA1000432	5.62	5.18	10.2	4.71	5.75	4.03	2.85	3.99	8.12		-	-
	Y79AA1000453	32.42	33.52	37.01	20.21	25.21	18.7	36.98	32.84	33.05	**	-	-
	Y79AA1000465	4.43	1.91	3.14	1.7	4.29	1.76	2.57	2.29	2.11		-	-
	Y79AA1000468	35.31	28.37	27.42	30.17	34.28	32.32	21.89	31.66	24.15		-	-
	Y79AA1000480	11.78	8.51	13.38	5.22	5.02	5.9	5.84	10.93	8.28	*	-	-
25	Y79AA1000502	19.24	16.15	21.16	8.66	14	12.35	16.39	18.08	18.17	*	-	-
	Y79AA1000521	75.16	71.09	84.73	74.85	78.28	67.85	48.04	51.73	45.89	**	-	-
	Y79AA1000534	13.31	15.15	20.4	13.25	14.05	10.71	13.15	11.71	11.75		-	-
	Y79AA1000538	12.36	14.67	20.45	9.58	11.27	5.96	3.97	0	4.64	**	-	-
	Y79AA1000539	15.51	16.32	14.99	9.48	12.74	10.1	5.39	5.2	6.13	* **	-	-
	Y79AA1000540	10.2	7.44	12.08	3.41	6.65	4.35	12.15	10.44	10.32	*	-	-
30	Y79AA1000560	180.34	128.09	149.28	162.08	194.06	150.66	118.75	125.1	108.26		-	-
	Y79AA1000574	12.15	9.72	10.83	5.66	8.8	4.53	10.59	16.21	10.56	*	-	-
	Y79AA1000584	40.28	35.5	37.1	23.89	28.68	25.51	26.96	34.59	37.41	**	-	-
	Y79AA1000589	9.65	4.15	9.45	2.79	3.53	2.1	3.54	8.38	4.6		-	-
	Y79AA1000598	34.26	29.42	37.32	17.43	15.95	16.46	24.99	27.55	21.09	** *	-	-
	Y79AA1000600	44.58	45.66	51.58	51.4	52.23	38.65	32.06	30.08	26.05	**	-	-
35	Y79AA1000609	7.91	6.61	8.76	1.95	5.66	3.3	3.74	4.36	7.05	*	-	-
	Y79AA1000618	20.27	20.13	21.51	6.61	13.31	6.56	19.86	14.92	16.1	**	-	-
	Y79AA1000627	18.35	16.53	18.66	9.09	10.58	9.21	14.78	8.21	18.69	**	-	-
	Y79AA1000636	5.21	2.52	7.03	3.36	4.46	4.2	2.54	3.13	1.94		-	-
	Y79AA1000649	19.51	21.35	25.26	12.58	12.75	9.69	16.99	19.63	19.13	**	-	-
	Y79AA1000656	7.37	5.14	7.48	2.86	2.97	2.28	2.83	3.72	2.99	** *	-	-
40	Y79AA1000673	8.38	7.17	11.58	4.74	3.99	5.74	1.87	4.27	2.18	* *	-	-
	Y79AA1000674	131.64	157.11	127.73	143.89	140.32	116.2	111.11	118.96	120.08		-	-
	Y79AA1000678	22.12	27.62	25.96	12.61	21.55	13.08	21.7	19.94	19.36	*	-	-
	Y79AA1000682	14.62	11.86	13.44	6.55	8.13	6.38	12.74	13.35	11.29	**	-	-
	Y79AA1000683	37.64	24.6	33.28	19.88	16.2	12.73	33.87	33.73	36.17	*	-	-
	Y79AA1000697	46.01	35.33	48.41	23.15	27.46	23.15	15.24	20.88	15.95	* **	-	-
45	Y79AA1000700	3.64	3.69	5.68	3.34	3.06	2.62	3.53	2.27	2.51		-	-
	Y79AA1000702	50.6	33.01	36	23.71	24.96	18.05	18.21	11.97	15.36	* *	-	-
	Y79AA1000704	93.86	71.31	88.5	110.92	93.16	106.68	74.22	61.46	46.95		-	-
	Y79AA1000705	6.23	6.74	7.37	1.82	2.22	2.09	2.28	0.26	2.39	** **	-	-
	Y79AA1000717	8.36	6.77	8.13	4.96	8.96	4.55	2.44	3.74	2.84	**	-	-
	Y79AA1000722	10.83	11.52	10.81	4.97	5.79	3.98	6.28	5.41	5.45	** **	-	-
50	Y79AA1000724	42.59	35.15	44.52	31.42	30.43	23.04	24.2	29.89	31.56	*	-	-
	Y79AA1000726	8.78	4.43	7.31	5.15	4.77	4.25	2.41	3.94	2.09	*	-	-
	Y79AA1000734	5.72	4.05	6.45	2.81	2.76	3.46	2.68	2.98	0.98	*	-	-
	Y79AA1000748	7.45	7.72	6.69	2.89	2.82	2.75	2.52	2.6	2.85	** **	-	-
	Y79AA1000750	41.52	33.1	33.65	18.03	22.65	18.6	26.15	29.63	22.6	** *	-	-
	Y79AA1000752	9.11	9.26	10.14	4.75	3.47	2.23	2.34	2.01	1.29	** **	-	-
55	Y79AA1000774	19.25	19.31	26.89	10.74	13.71	10.53	12.54	14.59	12.73	* *	-	-
	Y79AA1000776	12.05	9.97	9.69	5.12	6.81	2.36	10.58	10.24	7.92	*	-	-
	Y79AA1000777	17.61	11.59	13.15	7.32	8.15	7.21	6.02	5.12	6.68	* *	-	-

Table 504

	VESEN2000175	7.45	6.95	7.39	3.43	6.1	3.72	4.64	8.1	4.35	*	-	-
	VESEN2000186	18.04	20.73	21.8	11.19	14.77	10.58	8.74	12.84	12.61	**	**	-
5	VESEN2000199	11.71	6.48	9.96	8.59	9.24	5.52	6.89	6.26	14.79			
	VESEN2000200	150.21	142.83	144.63	172.94	215.31	128.05	93.38	87.48	97.05	**	**	-
	VESEN2000204	7.05	7.07	9.65	4.56	4.24	3	2.19	4.63	3.19	*	*	-
	VESEN2000218	16.65	12.07	17.29	7.31	8.24	6.16	8.32	5	5.15	**	**	-
	VESEN2000230	12.09	13.48	13.57	4.57	10.45	4.08	3.93	4.19	2.98	*	**	-
	VESEN2000272	8.95	9.01	11.14	4.32	3.66	3.22	4.36	3.53	4.22	**	**	-
10	VESEN2000299	8.61	7.23	11.81	5.04	5.83	4.25	3.6	5.71	4.76	*	*	-
	VESEN2000323	17.3	12.74	15.22	15.36	17.47	14.24	5.84	6.35	4.65	**	**	-
	VESEN2000327	18.5	9.19	14.45	6.81	8.17	7.45	3.2	5.07	5.85	*	*	-
	VESEN2000328	7.53	7.78	8.38	3.45	4.17	4.12	2.35	2.19	1.81	**	**	-
	VESEN2000330	25.07	20.15	22.73	13.34	18.45	13.16	15.63	15.15	15	*	**	-
	VESEN2000336	7.29	9.37	11.82	5.86	9.09	5.78	3.64	3.5	5	*	*	-
15	VESEN2000354	10.99	9.78	10.4	5.27	8.22	4.93	3.58	3.58	2.59	*	**	-
	VESEN2000378	168.58	110.88	139.59	81.44	98.69	74.23	74.17	76.59	54.75	*	*	-
	VESEN2000379	32.08	21.47	28.34	16.72	15.12	13	22.77	26.45	21.84	*	*	-
	VESEN2000397	21.17	10.11	25.4	12.77	12.2	10.8	15.72	16.01	17.99			-
	VESEN2000416	5.92	5.21	7.48	3.5	6.28	4.01	2.87	3.94	3.48	*	*	-
	VESEN2000420	8.47	4.35	7.18	4.04	3.81	2.46	2.31	4.27	1.92			-
20	VESEN2000430	8.74	5.76	7.33	2.94	5.32	2.11	1.8	2.27	0.71	*	**	-
	VESEN2000448	14.67	14.58	16.08	4.71	4.34	2.88	3.91	2.23	2.27	**	**	-
	VESEN2000449	46.1	50.29	39.97	32.75	32.82	21.56	37.62	28.99	23.73	*	*	-
	VESEN2000456	45.88	33.52	32.47	18.88	21.08	19.47	15.48	22.28	22.31	*	*	-
	VESEN2000562	6.11	3.13	4.69	5.4	3.95	3.19	3.24	2.45	2.42			-
	VESEN2000573	26.67	20.52	18.39	14.29	11.45	8.79	8.94	7.46	12.2	*	*	-
25	VESEN2000604	9.02	7.76	4.59	4.65	4.32	4.42	1.84	4.41	1.34	*	*	-
	VESEN2000614	9.59	7.78	6.05	4.77	5.05	4.35	3	3.94	2.87	*	*	-
	VESEN2000638	16.75	10.83	10.74	6.38	9.46	6.26	4.67	4.46	3.5	*	*	-
	VESEN2000641	15.71	14.29	14.25	7.95	7.6	6.71	8.94	7.41	6.6	**	**	-
	VESEN2000645	9.28	11.51	10.66	5.27	8.01	4.95	5.42	4.96	6.81	*	**	-
30	Y79AA1000013	24.66	15.88	15.09	6.08	5.43	5.53	13.59	13.07	18.88	*	*	-
	Y79AA1000030	19.52	16.62	14.47	8.07	7.77	10.99	7.86	7.77	10.02	*	**	-
	Y79AA1000033	29.91	18.65	28.38	16.37	15.72	11.59	28.15	19.64	28.85	*	*	-
	Y79AA1000037	4.84	2.8	4.89	5.26	3.24	0.91	1.44	1.94	2.43	*	*	-
	Y79AA1000041	79.98	53.74	54.67	39.77	36.14	29.21	42.77	30.09	45.97	*	*	-
	Y79AA1000059	9.8	7.3	8.44	4.37	5.08	2.06	5.32	3.01	5.11	*	*	-
35	Y79AA1000065	116.14	101.03	106.53	93.49	124.96	111.44	68.74	63.93	65.56	**	**	-
	Y79AA1000081	7.88	11.03	9.72	7.03	9.28	7.22	5.68	4.38	3.88	**	**	-
	Y79AA1000127	8	5.82	6.2	3.17	4.25	2.35	5.28	6.34	4.47	*	*	-
	Y79AA1000130	20.16	11.01	14.68	13.04	8.43	8.05	14.41	16.35	14.16			-
	Y79AA1000131	17.14	8.99	11.18	6.93	10.63	6.92	6.82	8.55	6.52			-
	Y79AA1000134	6.34	7.52	5.68	7.86	5.61	3.09	4.16	2.2	3.07	*	*	-
40	Y79AA1000143	8.89	7.38	8.14	3.93	5.83	3.75	3.23	2.53	3.96	*	**	-
	Y79AA1000144	11.79	8.62	9.18	5.64	6.37	5.56	3.97	3.84	3.36	*	**	-
	Y79AA1000150	28.57	28.05	20.04	12.4	17.77	10.86	24.46	16.09	20.29	*	*	-
	Y79AA1000153	11.32	11.62	7.41	5.89	9.36	6.53	6.84	5.59	6.52			-
	Y79AA1000166	13.75	6.3	9.45	4.16	5.68	2.88	7.42	8.85	8.76			-
	Y79AA1000179	13.15	8.67	7.74	7.91	7.5	4.67	13.52	5.81	9.18			-
45	Y79AA1000181	14.37	9.22	8.77	6.64	8.41	3.93	3.41	3.97	4.65	*	*	-
	Y79AA1000202	22.9	17.78	19.17	14.55	12.31	8.36	18.27	13.59	20.94	*	*	-
	Y79AA1000207	52.28	63.3	44.9	24.04	38.89	23.79	15.85	13.31	14.27	*	**	-
	Y79AA1000214	9.44	4.9	5.66	4.71	4.79	4.18	5.54	2.28	3.2			-
	Y79AA1000222	11.34	8.61	7.95	5.13	7.04	2.26	4.7	3.53	4.67	*	*	-
	Y79AA1000226	27.21	25.59	19.24	12.19	15.74	11.35	24.75	16.69	21.48	*	*	-
50	Y79AA1000227	9.07	6.57	6.62	3.66	4.45	3.9	7.4	6.45	7.51	*	*	-
	Y79AA1000230	32.97	25.47	29.26	12.55	10.05	6.68	4.18	3.82	4.93	**	**	-
	Y79AA1000231	65.27	41.2	48.87	47.4	47.89	24.64	28.97	32.14	28.29	*	*	-
	Y79AA1000239	74.35	85.68	67.35	68.07	52.75	45.41	64.49	54.97	44.91			-
	Y79AA1000258	8.42	5.48	5.2	4.36	4.01	3.09	3.34	2.93	4.62			-
	Y79AA1000268	80.53	75.19	49.36	22.18	30.03	27.42	30.39	28.87	31.5	*	*	-
55	Y79AA1000269	73.43	71.24	52.77	41.13	70.01	36.67	77.91	62.7	65.45			-
	Y79AA1000270	10.85	7.03	4.1	4.98	7.95	3.4	5.03	5.7	4.72			-
	Y79AA1000280	25.49	25.11	28.96	8.6	12.9	9.27	5.74	8.76	6.26	**	**	-

Table 503

	THYR01001730	7.74	3.57	7.41	4.48	4.19	3.07	3.64	4.09	4.72			
	THYR01001738	1.57	3.03	2.24	0.97	0.17	0.58	0.7	0.99	1.64	*	-	
5	THYR01001743	5.12	3.48	3.39	1.67	2.12	2.94	1.98	2.6	2.48			
	THYR01001745	30.25	14.84	20.6	6.81	4.65	9.25	12.7	8.09	11.05	*	-	
	THYR01001746	10.78	5.89	7.69	2.83	3.1	3.64	3.64	1.32	2.13	*	*	-
	THYR01001770	20.01	12.02	16.57	10.05	13	10.49	5.37	6.07	5.43	*	-	
	THYR01001772	29.2	56.08	29.11	28.47	34.34	15.95	9.21	8.24	9.37	*	-	
	THYR01001778	39.03	35.61	22.42	13.39	32.45	16.04	9.42	8.03	11.2	*	-	
10	THYR01001793	9.35	3.26	5.98	4.02	4.49	3.37	4.45	4.26	5.77			
	THYR01001796	66.96	62.34	52.36	31.61	31.58	30.06	74.83	63.67	78.59	**	-	
	THYR01001800	25.21	17.42	21.74	19.92	16.43	13.71	8.41	10.02	11.59	**	-	
	THYR01001803	12.96	11.11	10.61	7.04	8.3	3.44	5.71	5.48	6.64	*	**	-
	THYR01001809	10.44	7.96	6.8	4.59	3.72	3.71	3.62	1.76	1.34	*	**	-
	THYR01001817	18.78	13.28	15.51	15.03	13.53	6.93	8.79	4.88	8.37	*	*	-
15	THYR01001819	14.15	14.31	9.69	5.8	6.63	6.06	6.27	4.29	5.81	*	*	-
	THYR01001828	8.14	8.22	6.65	5.42	7.94	4.29	7.29	4.26	2.47			
	THYR01001854	50.53	32.88	37.71	33.17	34.18	22.81	22.16	27.13	21.15	*	-	
	THYR01001895	9.86	7.83	3.92	4.41	4.53	2.55	3.7	10.54	3.46			
	THYR01001907	27.03	13.41	12.49	9.48	10.54	7.02	4.92	4.41	6.47			
20	TRACH1000006	9.54	7.16	8.08	5.68	4.66	5.53	6.46	4.52	6.76	*	-	
	TRACH1000013	9.02	5.8	8.2	2.58	4.45	3.1	3.07	2.02	4.14	*	*	-
	TRACH1000074	13.69	9.32	8.63	4.66	5.76	4.06	6.37	3.07	3.63	*	*	-
	TRACH1000095	11.28	8.48	10.55	5.54	3.9	3.41	3.79	4.56	3.68	**	**	-
	TRACH1000102	26.86	26.3	20.1	15.81	18.57	15.2	32.52	24.6	27.09	*	-	
	TRACH1000108	88.66	59.61	40.48	28.57	26.39	20.8	69.3	56.08	66.91			
25	TRACH1000126	17.36	13.19	13.78	8.07	9.6	6.46	5.91	5.75	6.51	*	**	-
	TRACH1000146	7.25	6.83	5.07	5.29	3.75	3.17	3.49	3.25	4.97	*	-	
	TRACH1000160	11.76	11.24	7.7	4.05	5.59	4.95	5.32	4.22	5.87	*	*	-
	TRACH1000184	33.51	45.86	50.29	14.82	25.81	22.15	25.29	16.88	23.76	*	*	-
	VESEN1000004	11.35	8.9	8.13	5.46	9.59	6.32	5.53	5.44	6.01	*	-	
	VESEN1000007	32.64	30.13	26.81	16.7	21.99	18.54	37.52	29.73	33.89	**	-	
30	VESEN1000013	13.16	11.68	12.05	7.19	9.9	4.25	7.18	4.46	8.92	*	*	-
	VESEN1000028	23.17	20.53	28.94	10.16	15.29	14.76	9.13	12.72	8.48	*	**	-
	VESEN1000059	4.76	3.32	5.57	3.56	3.32	3.56	1.16	3.33	1.96			
	VESEN1000100	8.85	5.84	8.82	5.55	6.82	5.48	3.69	5.59	3.98	*	-	
	VESEN1000107	158.28	125.83	134.47	126.46	135.37	133.6	74.61	79.36	82.66	**	-	
	VESEN1000117	3.74	4.02	10.93	3.48	5.02	4.1	2.18	2.11	2.69			
35	VESEN1000122	5.84	5.31	7.17	3.91	7.47	3.29	2.52	3.75	2.41	*	-	
	VESEN1000137	7.93	10.91	7.7	8.31	5.46	4.02	3.34	3.36	5.77	*	-	
	VESEN1000195	12.47	4.75	10.07	4.43	4.22	7.07	3.11	4.32	13.25			
	VESEN1000215	89.55	69.15	78.83	74.46	93.71	81.06	50.08	36.95	43.24	**	-	
	VESEN1000279	16.42	11.12	19.97	7.86	19.23	9.52	8.84	13.86	21.33			
	VESEN1000363	40.43	29.46	56.48	41.4	53.72	50.13	17.51	33.82	20.25			
40	VESEN1000388	4.51	6.13	9.09	5.56	8.45	6.21	3.44	6.29	2.57			
	VESEN1000394	137.33	123.92	135.03	107.32	177.33	115.58	81.85	76.56	86.04	**	-	
	VESEN1000410	7.84	11.73	12.31	7.37	8.85	7.32	5.82	6.16	5.5	*	-	
	VESEN1000411	14.34	16.7	11.87	14.26	14.7	9.52	6.54	4.28	7.07	**	-	
	VESEN1000416	6.65	6.22	8.5	3.16	5.01	4.35	2.06	3.93	7.37	*	-	
	VESEN1000440	7.1	4.56	7.48	10.06	4.69	9.04	3.62	6.97	6.76			
45	VESEN1000452	127.08	114.54	102.58	100.33	106.23	122.79	77.87	58.16	82.71	*	-	
	VESEN1000539	57.4	47.68	67.32	36.3	45.7	42.57	51.56	51.15	50.18			
	VESEN1000554	7.29	12.24	14.13	6.05	4.57	6.75	3.69	3.4	3.85	*	-	
	VESEN1000557	4.5	6.91	10.86	5.64	5.73	4.35	3.05	2.96	4.75			
	VESEN1000575	16	18.27	24.28	11.3	16.07	11.77	13.59	13.83	16.27			
	VESEN1000585	14.4	14.28	13.38	9.87	13.37	6.72	4.91	4.08	5.01	**	-	
50	VESEN1000592	28.09	20.04	22.75	8.98	11.53	7	7.53	8.48	8.84	**	**	-
	VESEN1000658	9.27	5.98	6.99	4.92	3.93	3.11	3.31	12.18	6.06	*	-	
	VESEN1000669	7.65	11.62	18.09	8.2	9.11	7.91	4.21	6.95	3.61			
	VESEN1000743	27.12	25.27	28.49	19.46	23.17	19.77	12.01	15.14	12.53	*	**	-
	VESEN1000752	23.77	18.91	24.8	11.19	11.63	9.8	14.24	19.48	18.19	**	-	
	VESEN1000761	7.03	8.3	9.9	5.64	5.41	3.65	3.1	8.25	5.47	*	-	
55	VESEN2000039	16.58	20.4	16.46	7.12	12.23	7.45	5.23	4.66	6.91	*	**	-
	VESEN2000102	15.76	17.97	17.55	7.45	12.52	8.13	11.35	8.16	13.31	*	*	-
	VESEN2000164	69.66	54.85	54.07	18.32	20.39	15.6	15.69	21.51	20.97	**	**	-

Table 502

	THYR01001358	6.96	7.77	6.52	4.41	8.39	3.78	5.37	5.5	4.22	*	*	-
	THYR01001363	36.77	35.26	36.97	12.86	20.63	14.65	24.19	31.52	25.12	**	*	-
5	THYR01001365	2.39	1.31	4.42	2.3	2.69	1.43	1.82	2.18	1.5	*	**	-
	THYR01001374	10.11	11.14	11.66	5.19	7.75	3.64	2.84	5.3	4.44	*	*	-
	THYR01001401	11.45	8.59	8.77	5.02	5.3	5.96	5.5	8.56	6.31	*	*	-
	THYR01001403	3.12	3.62	7.98	1.61	2.43	2.61	3.73	1.98	3.91	*	*	-
	THYR01001405	84.82	60.76	92.35	68.75	78.6	50.19	53.53	29.6	47.94	*	*	-
	THYR01001406	13.59	7.57	12.02	4.76	6.87	3.76	11.69	13.59	9.54	*	*	-
10	THYR01001411	4.72	4.15	5.65	3.88	2.4	3.38	2.52	2.5	3.6	*	*	-
	THYR01001420	9.34	4.82	8.71	4.7	4.71	5.55	3.24	4.39	8.81	*	*	-
	THYR01001426	9.26	4.52	12.27	5.65	6.22	4.37	4.27	8.59	6.13	*	*	-
	THYR01001430	10.74	13.91	18.18	13.99	18.28	13.49	6.82	9.48	10.77	*	*	-
	THYR01001434	7.66	6.66	12.91	8.64	10.68	8.13	5.38	8.91	2.99	*	*	-
	THYR01001456	26.49	36.99	36.96	17.02	23.99	16.92	30.71	34.81	38.85	*	*	-
15	THYR01001457	14.33	10.91	13.26	5.72	4.54	3.33	4.05	3.22	4.83	**	**	-
	THYR01001458	15.32	13.83	12.78	10.99	10.13	5.72	4.4	7.29	6.97	*	**	-
	THYR01001459	5.78	4.92	7.36	2.81	3.46	4.24	3.11	4.43	5.51	*	*	-
	THYR01001471	13.48	10.81	13.3	5.13	9.62	6.19	6.36	8.85	9.45	*	*	-
	THYR01001478	8.04	3.79	8.01	2.94	3.62	2.21	4.12	4.22	3.59	*	*	-
	THYR01001480	16.07	16.86	14.48	6.78	13.68	13.34	3.54	6.61	19.35	*	*	-
20	THYR01001481	16.23	13.97	17.19	13.3	15.2	13.73	4.94	6.01	3.76	**	*	-
	THYR01001487	39.34	43.94	45.87	41.49	42.62	34.09	36.78	39.23	41.44	*	*	-
	THYR01001495	6.21	3.33	12.71	2.18	5.51	4.67	3.43	3.46	4.85	*	*	-
	THYR01001498	13.82	15.45	16.61	11.32	20.15	8.37	4.81	6.08	4.31	**	*	-
	THYR01001510	9.97	6.87	19.31	8.41	6.21	3.55	4.99	6.45	19.57	*	*	-
	THYR01001512	11.51	8.21	10.34	5.87	6.36	6.84	4.43	19.67	5.22	*	*	-
25	THYR01001519	12.05	16.47	10.9	4.73	8.69	4.14	4.42	4.28	5.04	*	**	-
	THYR01001522	10.39	6.84	12.83	6.15	7.71	8	5.13	5.33	7.56	*	*	-
	THYR01001523	15.14	10.51	16.01	7.01	9.2	6.38	3.77	3.19	4.5	*	**	-
	THYR01001526	5.67	5.48	12.84	2.85	5.19	3.09	4.51	3.63	4.01	*	*	-
	THYR01001529	70.52	99.68	61.07	73.62	81.82	35.88	29.07	23.9	34.69	*	*	-
	THYR01001534	10.48	7.92	11.72	6.09	7.42	11.24	6.53	6.35	3.17	*	*	-
30	THYR01001537	5.9	3.05	5.93	3.4	3.75	2.75	3.26	2.87	4.91	*	*	-
	THYR01001541	27.84	38.72	38.42	23.51	30.22	18.99	22.01	26.98	14.85	*	*	-
	THYR01001545	12.87	9.76	11.61	3.56	6.9	5.41	4.16	22.14	4.44	**	*	-
	THYR01001559	143.67	217.09	127.43	126.92	211.99	217.97	165.83	143.91	101.7	**	**	-
	THYR01001563	29.66	25.88	28.83	14.46	17.77	19.17	10.82	18.32	17.35	**	**	-
35	THYR01001570	5.85	5.24	10.72	3.21	3.79	2.31	2.47	4.03	3.05	*	*	-
	THYR01001573	10.58	14.52	15.67	7.66	10.98	10.26	4.93	9.17	5.76	*	*	-
	THYR01001584	11.71	9.79	13.05	7.75	12.43	7.3	8.19	4.92	4.88	*	*	-
	THYR01001593	119.63	87.93	122.57	37.27	54.3	33.56	100.89	84.41	119.61	**	*	-
	THYR01001595	9.11	9.13	10.8	7.84	8.88	6.77	4.62	12.97	7.82	*	*	-
	THYR01001596	6.25	3.54	8.97	3.03	4.5	2.76	3.54	3.67	4.02	*	*	-
40	THYR01001602	7.1	6.21	7.8	6.63	4.96	5.02	4.21	4.12	2.97	**	*	-
	THYR01001605	28.22	39.54	25.87	13.48	23.89	22.57	11.83	10.02	13.09	*	*	-
	THYR01001608	19.26	17.94	17.77	8.3	22	8.49	8.07	8.36	11.18	**	*	-
	THYR01001617	7.36	9.07	12.26	4.4	5.61	3.53	3.83	3.37	4.51	*	*	-
	THYR01001634	9.8	9.15	14.08	5.65	9.76	4.62	5.62	3.69	4.86	*	*	-
	THYR01001637	6.56	2.42	6.39	2.18	3.8	2.31	2.41	3.7	6.48	*	*	-
45	THYR01001641	12.87	8.05	12.44	6.03	4.87	6.49	3.32	2.86	2.92	*	**	-
	THYR01001656	20.03	12.66	30.89	7	10.07	5.22	6.32	7.53	5.64	*	*	-
	THYR01001658	7.58	6.85	10.14	4.65	4.77	3.94	3.38	2.8	2.5	*	**	-
	THYR01001661	150.1	98.3	106.51	135.25	166.2	125.8	74	80.32	53.76	**	**	-
	THYR01001671	10.97	8.43	12.21	5	5.04	3.97	5.12	5.12	3.84	**	**	-
	THYR01001672	12.9	11.31	11.75	5.85	6.91	5.08	3.69	2.74	2.72	**	**	-
50	THYR01001673	31.04	28.57	26.79	16.85	19.7	19.9	22.8	23.43	23.51	**	*	-
	THYR01001677	8.63	6.86	9.97	3.55	4.9	4.2	5.4	6.55	8.85	*	*	-
	THYR01001683	39.34	16.85	25.87	16.11	18.7	19.17	8.36	10.76	9.09	*	*	-
	THYR01001700	12.31	7.03	9.15	3.61	4.23	3.04	2.79	3.63	4.8	*	*	-
	THYR01001702	23.7	21.62	17.4	10.57	5.8	11.44	6.01	4.83	2.94	*	**	-
	THYR01001703	7.22	4.23	5.14	2.63	3.23	1.81	2.37	2.42	2.04	*	*	-
55	THYR01001706	8.39	5.62	8.09	3.72	3.11	3.37	3.24	2.65	1.26	*	**	-
	THYR01001721	11.69	10.42	16.73	6.09	4.96	4.46	3.92	2.33	3.87	*	**	-
	THYR01001725	147.69	144.4	144.74	111.99	128.3	86.63	130.43	106.61	93.76	*	*	-

Table S01

	THYRO1000934	9.21	6.94	9.67	4.17	7.52	4.26	6.41	5.73	4.96	*	*	-
	THYRO1000951	21.19	13.82	17.24	10.15	10.71	6.54	7.11	11.34	11.56	*	*	-
5	THYRO1000952	38.82	45.17	51.97	24.89	24.37	18.77	17.55	26.97	13.34	**	**	-
	THYRO1000956	11.42	8.41	13.85	5.27	6.34	6.15	5.07	6.61	6.85	*	*	-
	THYRO1000960	9.97	3.83	8.63	2.78	4.13	2.34	2.46	2.03	6.43	*	*	-
	THYRO1000961	33.04	35.26	36.48	15.01	15.48	15.21	21.6	24.41	21.14	**	**	-
	THYRO1000964	14.43	12.46	17.93	3.92	6.7	5.62	5.43	7.53	4.58	**	**	-
	THYRO1000971	15.18	16.5	16.14	8.34	13.54	9	9.38	8.59	10.52	*	**	-
10	THYRO1000974	7.51	5.29	9.43	2.24	6.94	1.8	3.55	5.59	3.45			-
	THYRO1000975	7.71	4.75	6.93	3.36	5.71	4.79	3.77	4.87	3			-
	THYRO1000983	8.4	4.14	7.44	3.86	5.03	3.36	3.1	3.21	4.03			-
	THYRO1000984	9.35	8.4	11.52	6.49	7.32	5.22	5.08	2.55	2.65	*	**	-
	THYRO1000988	10.2	6.57	7.62	4.27	4.23	4.6	2.94	2.66	2.92	*	**	-
	THYRO1000991	12.33	10.95	12.47	5.27	7.49	7.34	5.57	6.15	6.01	**	**	-
15	THYRO1000999	9.77	12.84	10.64	5.68	8.33	5.92	5.68	3.11	2.9	*	**	-
	THYRO1001003	324.55	443.5	324.63	353.93	491.59	377.12	509.02	394.65	380.17			-
	THYRO1001015	16.58	13.37	11.66	7.36	13.81	8.4	12.04	9.04	10.47			-
	THYRO1001016	34.7	29.76	32.31	14.48	17.7	10.51	27.97	31.88	30.58	**		-
	THYRO1001022	27.6	17.2	20.61	6.98	9.39	10.85	5.35	6.58	6.8	*	**	-
20	THYRO1001031	15.75	9.57	11.29	5.12	7.05	5.99	8.13	6.39	9.34	*		-
	THYRO1001033	11.34	7.57	8.58	7.55	5.52	6.01	4.3	4.01	4.9	*	*	-
	THYRO1001062	12.86	12.45	12.49	4.68	5.59	4.88	3.54	3.68	2.79	**	**	-
	THYRO1001063	10.18	8.38	10.08	5.14	7.17	4.58	3.55	4.32	2.35	*	**	-
	THYRO1001071	18.49	15.5	18.23	9.12	10.91	7.86	5.73	5.41	5.84	**	**	-
	THYRO1001080	11.77	12.06	11.47	3.65	12.27	3.17	3.08	4.59	4.41	**		-
25	THYRO1001093	12.17	9.63	9.75	5.98	4.91	5	8.58	8.31	7.24	**		-
	THYRO1001100	33.06	27.59	26.02	30.29	27.98	30.28	66.02	40.22	70.38	*		+
	THYRO1001102	10.18	6.7	5.95	5.04	6.9	4.52	4.36	4.71	4.45			-
	THYRO1001104	5.1	2.75	2.78	0.69	0.92	1.62	1.32	0.94	0.7	*	*	-
	THYRO1001109	19.62	18.07	14.26	10.98	10.98	9.76	8.55	11.06	8.68	*	*	-
	THYRO1001113	10.89	6.97	7.59	2.84	3.69	2.77	2.56	1.4	2.3	*	**	-
30	THYRO1001120	9.27	9	8.82	3.46	4.66	2.98	4.02	3.07	2	**	**	-
	THYRO1001121	20.41	16.36	16.91	9.1	13.05	6.93	13.22	15.3	13.97	*		-
	THYRO1001128	25.09	18.24	17.74	8.53	9.61	6.74	3.45	4.7	7.82	**	**	-
	THYRO1001133	7.42	6.4	5.88	4	5.05	4.23	3.88	4.14	3.81	*	**	-
	THYRO1001134	29.23	23.66	23.28	16.51	20.53	18.49	22.14	16.92	20.47	*		-
	THYRO1001142	5.04	4.99	5.1	2.78	3.05	3.47	2.44	2.26	1.6	**	**	-
35	THYRO1001173	31.85	26.04	28.94	20.36	18.72	13.59	18.5	15.33	18.17	*	**	-
	THYRO1001175	95.55	98.04	80.56	104.29	114.18	77.1	51.33	46.56	49.17	**		-
	THYRO1001177	10.9	11.18	11.71	5.13	7.17	5.9	4.63	5.82	5.02	**	**	-
	THYRO1001189	16.52	14.55	11.53	5.76	11.24	3.42	11.07	10.9	8.86			-
	THYRO1001194	11.83	6.2	11.19	5.29	7.22	4.28	5.84	7.01	6.86			-
	THYRO1001204	52.04	33.32	32.8	18.53	22.19	18.27	39.75	37.94	46.59	*		-
40	THYRO1001205	8.98	12.23	9.39	5.39	8.08	5.04	6.55	6.04	4.85	*	*	-
	THYRO1001213	41.54	37.73	35.86	23.36	25.7	20.79	37.67	32.41	38.62	**		-
	THYRO1001224	10.86	11.43	10.75	5.38	8.69	6.7	7.21	4.53	6.36	*	**	-
	THYRO1001237	16.76	20.25	15.97	16.59	13.69	5.38	9.78	5.69	6.27	**		-
	THYRO1001242	13.1	19.03	16.06	12.27	8.61	6.73	9.21	5.95	9.02	*	*	-
	THYRO1001258	27.32	21.68	20.65	9.74	18.76	12.77	10.51	6.86	13.04	*	**	-
45	THYRO1001262	8.57	4.74	4.06	3.76	5.55	4.02	4.79	3.93	5.05			-
	THYRO1001266	25.46	17.38	17.48	9.96	11.28	8.76	13.97	15.24	19.55	*		-
	THYRO1001271	27.85	25.6	17.14	13.77	18.43	9.74	22.54	18.92	23.05			-
	THYRO1001287	126.74	172.63	126.71	56.76	169.06	71.99	103.16	93.81	120.13			-
	THYRO1001290	11.3	8.82	9.68	4.97	7.48	6.55	3.86	5.84	8.41	*		-
	THYRO1001291	13.9	16.44	13.22	8.85	8.33	3.9	6.83	3.64	6.66	*	**	-
50	THYRO1001297	13.17	12.27	11.41	6.73	6.89	4.23	7.87	5.3	8.04	**	**	-
	THYRO1001302	28.88	24.47	17.01	9.99	19.11	8.25	41.39	28.42	36.98			-
	THYRO1001313	3.4	2.28	2.86	1.62	1.97	1.71	3.09	2.85	2.3	*		-
	THYRO1001320	4.85	2.3	4.45	3.81	2.95	2.21	2.67	2.22	1.94			-
	THYRO1001321	84.54	108.43	89.51	45.59	48.9	36.67	60.61	56.13	42.33	**	*	-
	THYRO1001322	15.07	13.59	11.8	5.6	6.9	4.45	5.91	6.73	6.71	**	**	-
55	THYRO1001327	61.14	62.79	47.56	27.21	36.08	29.25	40.32	44.89	30.36	**	*	-
	THYRO1001336	21.04	18.95	13.64	17.33	15.44	11.88	17.21	14.24	12.35			-
	THYRO1001347	42.47	35.49	29.56	20.44	37.43	12.7	20.26	16.34	15.93	**		-

Table 500

	THYRO1000558	8.54	7.17	6.38	4.15	8.61	3.01	4.26	2.04	3.45	*	-
	THYRO1000569	6.63	4.92	4.79	4.77	4	3.41	3.56	3.02	3.94	*	-
5	THYRO1000570	9.09	6.95	8.33	5.27	6.38	6.54	5.25	4.89	7.09		
	THYRO1000572	52.16	31.52	39.78	12.92	33.15	30.73	30.68	23.84	11.23		
	THYRO1000573	8.96	8.49	6.82	4.33	8.49	6.35	13.45	11.85	8.38		
	THYRO1000577	9.73	7.83	7.69	3.93	6.26	2.6	2.58	3.82	2.74	*	**
	THYRO1000580	40.56	32.54	29.95	22.22	19.86	19.81	24.19	17.34	25.52	*	*
	THYRO1000584	39.96	36.8	31.93	19.25	25.5	16.24	26.15	21.02	19.13	*	*
10	THYRO1000585	8.99	7.37	6.65	4.19	6.67	4.86	5.4	3.58	2.99	*	-
	THYRO1000596	43.85	33.14	32.61	22.94	31.65	11.59	15.52	10.88	14.47	**	-
	THYRO1000602	19.04	15.25	10.78	7.23	11.51	11.78	8.87	7.72	10.81		
	THYRO1000605	10.06	9.44	5.91	6.29	6.99	4.61	4.9	5.21	5.08		
	THYRO1000615	134.15	117.37	85.97	86.81	98.31	67.18	77.31	67	84.37		
	THYRO1000625	33.27	24.69	16.68	12.6	10.75	20.13	8.38	6.86	11.76	*	-
15	THYRO1000636	17.91	16.49	14.48	12.48	11.49	6.7	8.72	6.17	8.59	*	**
	THYRO1000637	18.22	22.2	13.43	10.39	14.05	9.78	9.07	5.2	7.26	*	-
	THYRO1000641	7.23	5.87	6.32	4.82	10.06	3.87	8.42	4.55	2.07		
	THYRO1000657	10.91	9.86	8.65	6.93	7.74	7.68	9.53	8.35	10.94	*	-
	THYRO1000658	14.37	9.48	5.1	6.28	6.83	5.43	4.59	4.62	4.77		
	THYRO1000662	10.63	9.43	8.46	6.7	6.85	6.96	7.92	7.09	10.89	*	-
20	THYRO1000666	30.51	33.5	45.43	12.9	20.33	13.81	14.6	13.53	16.48	*	**
	THYRO1000676	11.56	11.35	8.48	6.32	9.3	8.37	4.16	5.02	6.44	*	-
	THYRO1000678	23.44	18.73	20.73	10.13	15.99	8.49	9	7.38	9.87	*	**
	THYRO1000684	27.78	27.85	20.27	12.26	16.45	11.06	26.97	17.92	24.98	*	-
	THYRO1000694	16.87	11.78	10.72	5.47	10.25	5.99	12.29	9.36	8.6		
	THYRO1000699	9.11	6.36	4.55	5.62	6.31	3.48	7.37	3.08	5.06		
25	THYRO1000712	17.55	14.87	12.31	9.08	11.99	7.67	12.79	9.26	13		
	THYRO1000715	29.82	23.25	21.02	20.98	27.37	20.44	15.76	18.98	18.68		
	THYRO1000716	13.12	15.14	11.94	8.87	12	7.85	7.48	6.69	6.88	**	-
	THYRO1000717	11.89	15.21	12.45	6.9	9.22	5.08	30.93	6.96	7.18	*	-
	THYRO1000723	30.36	26.4	21.78	10.82	20.63	14.48	30.94	22.3	30.26	*	-
30	THYRO1000734	16.61	19.91	16.12	11.94	17.61	9.39	9.31	6.61	13.3	*	-
	THYRO1000748	17.46	14.06	15.87	5	7.52	5.42	7.82	5.84	7.09	**	**
	THYRO1000755	20.17	21.24	19.88	13.28	17.61	14.04	14.7	23.04	21.01	*	-
	THYRO1000756	369.81	334.57	369.63	298.51	345.41	273.28	259.37	366.07	273.4		
	THYRO1000776	4.97	4.99	7.55	4.25	6.18	4.82	2.58	5.02	3.55		
	THYRO1000777	5.42	5.63	6.34	5.34	6.01	4.03	3.82	3.75	3.21	**	-
35	THYRO1000779	483.37	458.9	483.51	453.87	549.42	449.1	368.6	510.93	458.47		
	THYRO1000782	17.27	20.25	23.35	14.58	17.12	13.37	15.61	16.4	16.35		
	THYRO1000783	5.74	5.45	6.68	5.43	9.82	4.88	3.62	5.19	3.8	*	-
	THYRO1000786	25.62	30.48	26.26	16.78	13.18	14.48	32.89	30.78	26.94	**	-
	THYRO1000787	50.52	37.21	54.05	28.71	18.06	26.25	20.11	24.5	22.66	*	**
	THYRO1000792	14.11	12.1	15.65	4.55	6.72	7.1	8.13	7.14	8.33	**	**
40	THYRO1000793	36.92	35.83	42.61	19.08	27.16	19.55	34.91	38.74	32.64	**	-
	THYRO1000795	37.33	45.29	88.32	42.19	57.36	40.15	12.57	21.17	13.38		
	THYRO1000796	10.25	11.83	14.01	6.19	10.21	5.98	7	11.89	11.1		
	THYRO1000798	9.87	14.22	12.83	9.06	8.1	7.49	5.04	7.33	5.2	*	*
	THYRO1000800	37.69	54.75	19.73	33.2	22.38	23.21	13.79	16.89	13.75		
	THYRO1000805	8.27	6.21	6.02	5.33	8.39	4.44	3.81	4.14	3.82	*	-
45	THYRO1000815	109.41	92.13	90.62	43.51	38.14	25.61	31.76	21.16	31.98	**	**
	THYRO1000829	33.86	25.13	34.71	19.68	21.68	18.45	16.87	14.34	12.65	*	**
	THYRO1000835	9.51	9.82	10.3	6.71	6.28	4.65	5.41	5.24	6.07	**	**
	THYRO1000843	27.06	16.5	24.96	16.28	19.34	15.73	6.22	7.81	8.06	**	-
	THYRO1000846	7.86	5.91	13.48	4.53	6.28	7.74	3.89	7.02	5.26		
	THYRO1000852	24.69	25.37	23.66	12.87	20.2	12.5	9.33	10.39	11.62	*	**
50	THYRO1000855	7.12	7.47	10.43	2.52	5.79	2.88	4.3	6.1	5.2	*	-
	THYRO1000865	9.21	5.67	8.64	3.4	6.56	3.18	5.99	8.76	7.67		
	THYRO1000866	57.27	46.1	51.69	32.85	43.74	27.97	46.24	39.05	41.55	*	-
	THYRO1000881	13.67	12.21	13.18	9.6	10.62	12.24	10.21	16.82	10.91		
	THYRO1000894	9.33	7.11	11.4	5.66	6.12	4.31	4.14	4.68	4.75	*	*
	THYRO1000895	8.9	5.89	10.11	3.35	4.07	4.72	4.32	4.09	3.45	*	*
55	THYRO1000916	9.19	7.72	16.87	5.02	6.67	5.27	4.16	6.12	3.42		
	THYRO1000917	6.89	10.63	12.32	4.97	9.51	5.39	6.77	7.94	6.2		
	THYRO1000926	14.99	10.81	12.37	6.79	9.95	6.06	7.42	7.03	11.27	*	-

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	THYRO1000087	51.14	69.3	41.47	29.99	58.4	26.89	61.05	53.08	61.77				
	THYRO1000092	25.07	28.19	30.84	13.75	13.51	9.94	17.56	21.51	22.34	**	*	-	-
5	THYRO1000093	4.26	2.01	4.16	1.77	3.46	3.25	2.69	3.3	2.41				
	THYRO1000099	9.62	5.87	9.26	3.68	8.63	4.69	5.9	7.16	4.32				
	THYRO1000107	70.48	59.74	59.68	39.34	64.57	40.78	35.76	61.82	50.34				
	THYRO1000111	13.96	11.92	12.38	7.04	9.78	5.89	9.55	11.53	9.66	*	*	-	-
	THYRO1000121	31.45	32.43	38.13	15.06	18.17	17.71	26.05	28.32	34.79	**		-	-
	THYRO1000124	54.83	35.35	48.22	37.6	35.37	31.52	34.87	27.99	31.75				
10	THYRO1000129	8.59	9.01	8.49	5.14	9.82	6.06	8.93	8.73	10.55				
	THYRO1000130	12.23	8.25	10.08	6.62	6.95	6.32	6.99	9.76	9.68	*		-	-
	THYRO1000132	7.93	8.79	12	5.35	7.15	7.04	4.12	4.66	3.86	*		-	-
	THYRO1000134	17.2	6.06	9.3	6.24	6.27	12.5	5.33	7.31	5.41				
	THYRO1000144	23.56	30.56	32.27	27.55	32.97	28.83	18.04	19.4	19.02	*		-	-
	THYRO1000155	35.26	37.35	35.3	18.99	24.31	20.42	31.31	40.6	29.86	**		-	-
15	THYRO1000156	37.13	43.79	38.99	24.77	34.58	23.66	18.78	36.39	21.81	*		-	-
	THYRO1000163	56.18	53.4	58.15	23.96	28.08	23.46	11.7	14.01	16.02	**	**	-	-
	THYRO1000173	8.52	8.35	6.58	5.25	10.18	4.15	3.25	4.22	4.73	**	**	-	-
	THYRO1000186	35.39	31.09	36.48	14.41	13.8	14.41	7.62	12.05	15.91	**	**	-	-
	THYRO1000187	17.39	15.95	24.02	6.63	8.73	8.87	5.49	7.02	6.9	*	**	-	-
	THYRO1000190	7.06	7.97	10.31	6.43	5.1	6.9	4.78	5.24	5.17	*		-	-
20	THYRO1000196	8.72	14.15	13.33	14.25	10.93	7.74	3.75	6.65	9.84				
	THYRO1000197	13.75	17.13	15.02	14.96	17.34	16.5	8.34	11.47	6.36	*		-	-
	THYRO1000199	6.8	9.71	11.38	6.6	11.32	10.32	5.5	6.51	7.17				
	THYRO1000206	20.09	21.74	25.04	13.14	17.5	12.03	7.45	6.8	7.63	*	**	-	-
	THYRO1000221	6.96	7.15	5.44	4.48	8.31	3.94	3.81	4.45	3.66	*		-	-
	THYRO1000222	56.33	56.17	63.27	37.58	40.7	31.29	35.13	33.5	43.65	**	**	-	-
25	THYRO1000228	148.64	127.09	184.25	73.36	64.55	57.59	55.06	55.21	55.09	**	**	-	-
	THYRO1000241	7.79	5.72	9.15	4.42	5	4.49	2.68	4.65	4.26	*	*	-	-
	THYRO1000242	18.16	23.99	29.76	25.09	51.85	27.45	12.5	20.02	16.61				
	THYRO1000246	21.57	18.12	24.56	19.46	20.13	15.33	9.29	10.17	9.72	**		-	-
	THYRO1000253	10.77	13.3	16.4	9.6	16.01	8.45	6.71	8.04	5.73	*		-	-
	THYRO1000270	12.58	12.99	17.24	8.11	13.02	8.87	9.06	8.16	7.99	*		-	-
30	THYRO1000279	15.04	13.53	16	8.28	14.2	7.73	7.71	7.11	4.92	**		-	-
	THYRO1000285	1215.2	1221.3	1148.5	961.64	1121.8	816.57	1261.8	1135.7	1518.7				
	THYRO1000288	24.11	22.65	22.01	13.89	25.53	14.41	22.83	21.93	20.34				
	THYRO1000296	40.86	36.2	38.51	33.51	37.84	29.24	32.78	49.02	40.29				
	THYRO1000320	12.29	9.4	14.15	6.59	7.17	6.08	5.28	11.64	9.29	*		-	-
35	THYRO1000322	37.49	34.76	35.08	17.69	26.05	19.02	19.43	23.34	20.74	**	**	-	-
	THYRO1000327	29.25	26.35	19.48	16.99	27.29	15.75	10.07	12.24	13.52	*		-	-
	THYRO1000343	16.65	12.58	17.82	8.18	11.42	7.21	10.1	9.05	8.44	*	*	-	-
	THYRO1000345	6.25	7.77	6.06	4.74	8.59	3.02	4.68	4.72	4.12	*	*	-	-
	THYRO1000358	13.06	10.33	11.35	6.24	8.76	7.32	8.15	6.51	9.21	*	*	-	-
	THYRO1000368	5.38	5.26	6.26	3.15	3.94	2.68	6.19	3.06	2.43	**		-	-
40	THYRO1000375	14.1	11.6	18.84	8.56	5.99	5.1	3.89	3.2	3.55	*	**	-	-
	THYRO1000381	10.31	8.3	9.78	4.37	5.3	5.95	4.17	5.35	3.59	**	**	-	-
	THYRO1000387	11.39	10.61	13.54	4.54	5.97	3.46	10.59	5.84	3.07	**		-	-
	THYRO1000394	61.78	50.97	22.02	14.67	26.91	8.08	6.98	9.73	11.33	*		-	-
	THYRO1000395	29.06	37.87	27.54	18.13	24.45	14.15	25.79	23.19	25.7	*		-	-
	THYRO1000400	10.51	10.51	9.69	3.23	12.01	5.83	3.72	3.73	3.08	**	**	-	-
45	THYRO1000401	10.42	9.53	10.77	6.74	5.06	4.73	3.24	5.18	5.6	**	**	-	-
	THYRO1000407	111.95	92.99	98.17	63.29	75.05	58.17	51.33	55.33	55.02	**	**	-	-
	THYRO1000420	15.01	13.25	17.92	11.1	11.62	8.56	12.79	12.37	14.41	*		-	-
	THYRO1000438	8.38	5.22	8.85	5.12	4.11	4.31	4.22	4.7	3.13	*	*	-	-
	THYRO1000452	15.04	12.59	12.97	5.92	10.6	6.13	2.16	3.93	4.57	*	**	-	-
	THYRO1000455	7.17	8.57	9.52	2.62	3.31	3.44	2.83	2.18	2.11	**	**	-	-
50	THYRO1000471	78.16	81.99	62.01	65.6	82.48	70.86	81.77	51.28	74.06				
	THYRO1000481	8.9	7.89	8.46	4.91	7.54	4.12	2.78	3.16	5.02	**		-	-
	THYRO1000484	19.83	12.76	16.87	10.37	11.59	9.38	11.21	9.7	11.91	*		-	-
	THYRO1000488	28.64	20.75	23.35	20.89	21.13	19.59	15.6	20.62	19.42				
	THYRO1000501	11.59	12.72	13.28	7.74	9.03	6.79	8.05	7.95	10.26	**	*	-	-
	THYRO1000502	55.86	52.65	49.31	44.07	46.78	38.57	44.58	40.93	38.11	*	*	-	-
55	THYRO1000505	9.64	7.65	7.27	5.56	5.82	3.06	3.98	5.47	2.02	*	*	-	-
	THYRO1000535	37.32	41.34	38.41	38.72	38.24	31.02	23.49	20.01	18.32	**		-	-
	THYRO1000556	102.92	92.57	85.3	76.31	104.96	62.73	51.56	50.77	49.93	**		-	-

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	SKHNC1000104	16.76	17.55	17.05	8.52	14.24	10.52	9.72	10.59	13.24	*	**	-	-
	SKHNC1000113	45.53	34.91	35.51	20.7	25.14	23.85	18.86	24.14	23.6	*	*	-	-
5	SKHNC1000119	17.58	14.67	21.54	10.33	12.31	8.53	8.83	13.02	8.17	*	*	-	-
	SKHNC1000142	16.52	20.57	23.49	7.26	16.72	11.75	8.01	11.68	14.09	*	*	-	-
	SKHNC1000170	242.49	183.41	211.38	201.28	226.18	182.82	247.45	216.7	250.73			-	-
	SKHNC1000178	15.01	12.68	14.32	6.9	9.12	6.49	4.97	7.82	9.01	**	**	-	-
	SKHNC1000194	7.33	8.61	11.61	2.23	3.73	3.19	2.64	4.74	3.45	*	*	-	-
	SKHNC1000198	18.76	29.87	24.52	7.87	13.1	9.06	5.41	5.23	5.02	*	**	-	-
10	SKHNC1000225	100.44	119.68	87.32	52.94	84.28	50.45	35.91	33.78	44.06		**	-	-
	SKHNC1000249	23.35	22.68	30.9	13.07	19.76	17.18	12.83	14.11	14.19	*	*	-	-
	SPLN1000007	10.7	10.7	16.74	6.92	9.27	7.23	6.86	7.37	5.89	*	*	-	-
	SPLN1000012	15.31	13.75	18.53	8.26	13.65	10.56	13.22	11.71	11.02			-	-
	SPLN1000014	74.22	59.99	63.53	32.33	49.13	32.58	29.87	29.67	37.64	*	**	-	-
	SPLN1000036	14.69	11.54	10.05	7.95	8.19	6.93	3.18	3.76	2.42	*	**	-	-
15	SPLN1000059	39.71	33.5	32.87	12.1	19.81	12.24	19.97	18.35	17.18	**	**	-	-
	SPLN1000068	20.63	22.57	21.07	11.6	16.29	10.46	9.19	11.89	8.79	*	**	-	-
	SPLN1000072	19.88	16.93	14.96	6.06	9.86	7.47	5.18	4.49	6.78	**	**	-	-
	SPLN1000101	71.97	43.16	57.46	18.69	27.56	26.8	32.73	40.27	19.97	*	*	-	-
	SPLN1000108	7.69	6.28	7.07	3.65	4.35	2.67	3.12	2.84	2.04	**	**	-	-
	SPLN1000113	11.47	6.64	12.13	7.54	8.23	7.26	5.5	6.27	7.59			-	-
20	SPLN1000114	16.01	14.75	14.36	7.29	9.65	6.3	8.14	8.22	8.73	**	**	-	-
	SPLN1000132	5.96	4.51	7.38	2.74	4.42	2.34	1.47	2.25	2.21	*	*	-	-
	SPLN1000135	10.57	8.18	9.39	3.86	4.71	4.73	4.9	2.38	2.97	**	**	-	-
	SPLN1000136	23.91	23.93	20.23	10.77	11.86	10.15	16.69	14.04	13.77	**	**	-	-
	SPLN1000141	42.88	39.73	42.83	27.16	32.91	23.4	23.31	18.41	20.57	**	**	-	-
	SPLN1000164	15.72	13.33	13.69	7.76	12.46	6.89	9.59	8.11	9.12	**	*	-	-
25	SPLN1000166	12.98	9.08	11.45	4.56	6.01	5.04	5.49	5.17	6.32	**	*	-	-
	SPLN1000175	19.26	16.94	15.5	7.48	9.23	6.92	11.34	13.49	13.41	**	*	-	-
	SPLN1000182	67.34	58.35	68.22	30.69	30.43	27.88	28.16	29.1	23.7	**	**	-	-
	SPLN1000185	6.38	8.35	5.56	4.69	4.56	3.06	2.86	2.31	1.51	**	*	-	-
	THYMU1000004	48.79	34.96	41.97	30.97	32.65	27.64	26.98	17.5	13.12	*	*	-	-
30	THYMU1000009	14.59	13.55	14.88	8.47	10.1	5	7.34	4.91	5.62	*	**	-	-
	THYMU1000015	19.34	18.55	12.08	8.34	11.37	5.93	10.39	10.96	13.03	*	*	-	-
	THYMU1000016	6.78	4.3	5.54	2.79	2.51	1.91	5.14	3.5	4.34	*	*	-	-
	THYMU1000023	6.35	6.23	6.53	4.39	3.65	4.2	3.1	5.69	3.98	**	*	-	-
	THYMU1000034	390.65	392.5	375.97	306.71	379.09	270.69	454.49	361.08	369.56			-	-
	THYMU1000035	8.93	7.8	9.14	5.88	4.95	4.81	5.06	3.93	4.36	**	**	-	-
35	THYMU1000037	9.69	6.41	7.36	4.28	3.96	3.33	4.24	3.59	3.83	*	*	-	-
	THYMU1000042	6.85	3.93	5.82	2.75	3.24	2.02	2.08	1.63	1.09	*	*	-	-
	THYMU1000047	25.64	25.03	24.74	13.43	16.94	11.38	26.77	21.68	23.81	**	*	-	-
	THYMU1000080	20.17	51.54	27.22	17.42	19.98	26.99	13.06	15.8	14.07			-	-
	THYMU1000094	11.12	6.99	9.51	6.45	6.49	4.35	11.14	9.41	9.31			-	-
	THYMU1000109	228.62	201.78	195.22	195.65	173.29	154.39	235.81	205.84	221.79			-	-
40	THYMU1000127	44.2	30.97	37.5	12.05	18.11	17.75	24.49	15.04	18.2	**	*	-	-
	THYMU1000130	21.64	17.3	15.84	7.65	12.45	8	10.24	3.41	8.22	*	*	-	-
	THYMU1000137	6.81	10.03	8.23	3.35	5.2	3.34	4.41	2.7	3.47	*	*	-	-
	THYMU1000146	11.97	7.39	7.48	8.11	7.24	6.17	5.13	3.03	6.19			-	-
	THYMU1000159	10.45	10.21	8.17	6.28	6.42	3.77	4.54	3.62	4.78	*	**	-	-
	THYMU1000163	8.08	8.93	9.7	5.33	5.95	3.66	7.57	4.17	4.21	**	*	-	-
45	THYMU1000167	67.79	51.53	54.02	27.07	31.06	17.27	75.27	48.55	70.31	**	*	-	-
	THYMU1000186	8.38	8.15	3.89	3.94	4.49	2.65	3.36	2.91	3.02			-	-
	THYR01000017	24	17.58	11.71	8.21	11.63	5.25	6.04	5.8	4.42	*	*	-	-
	THYR01000026	10.24	8.59	6.72	5.28	7.77	4.55	4.15	3.45	6.46	*	*	-	-
	THYR01000034	39.83	32.92	35.9	29.82	38.02	28.83	26.88	25.91	22.08	*	*	-	-
	THYR01000035	34.67	27.15	24.29	22.39	30.44	22.35	22.09	19.28	23.65			-	-
50	THYR01000036	16.2	14.28	11.79	9.92	10.8	4.36	7.94	4.75	8.23	*	*	-	-
	THYR01000040	9.47	7.71	9.48	4.32	5.83	3	5.07	3.54	5.01	*	**	-	-
	THYR01000061	18.28	19.76	15.63	8.28	5.85	4.81	6.42	5.28	9.19	**	**	-	-
	THYR01000067	16.85	13.02	12.06	8.02	8.96	4.34	7.22	6.1	5.44	*	**	-	-
	THYR01000070	9.96	13.03	10.58	5.63	9.48	4.69	20.02	18.33	15.63	*	*	-	-
	THYR01000072	21.57	19.92	18.73	19.89	17.94	12.39	9.77	9.74	9.5	**	*	-	-
55	THYR01000084	9.91	5.58	5.31	3.9	6.43	2.97	10.41	8.45	10.27			-	-
	THYR01000085	29.04	36.56	22.76	13.84	17.05	13.37	10.59	14.52	10.12	*	*	-	-
	THYR01000086	13.56	10.42	8.08	7.5	9.36	4.87	6.64	6.5	4.47			-	-

Table 497

	PLACE4000392	8.31	9.76	11.03	3.63	5.95	3.64	5.73	2.01	2.93	**	*	-	-
	PLACE4000399	53.9	51.54	53.26	20.64	28.55	18.63	37.94	35.1	44.54	**	**	-	-
5	PLACE4000401	4.22	3.76	3.44	3.96	3.49	2.6	3.5	3.9	1.61			-	-
	PLACE4000403	8.42	5.82	6.5	4.19	7.67	4.85	3.96	3.29	3.53	*		-	-
	PLACE4000411	20.23	15.5	15.73	8.23	11.19	9.81	14.98	10.1	11.34	*		-	-
	PLACE4000415	8.15	4.8	4.61	4.06	6.07	4.23	2.98	3.98	4.5			-	-
	PLACE4000416	35.43	25.11	24.69	17.72	17.34	16.23	24.55	17.6	25.73	*		-	-
	PLACE4000424	43.93	33.38	31.83	13.69	19.37	13.98	27.93	23.64	33	**		-	-
10	PLACE4000431	10.44	8.78	8.78	5.88	4.85	3.5	4.69	2.33	4.08	**	**	-	-
	PLACE4000443	50.64	44.47	41.43	26.49	37.08	21.33	51.22	33.11	43.74	*		-	-
	PLACE4000445	5.42	4.17	3.6	3.57	7.43	5.38	2.23	3.95	2.66			-	-
	PLACE4000450	9.85	6.68	6.23	4.72	6.64	4.07	3.56	7.58	3.73			-	-
	PLACE4000455	167.91	132.51	128.81	98.87	124.22	95.79	172.16	139.08	172.24			-	-
	PLACE4000465	31.84	25.74	18.83	10.79	23.42	11.27	11.29	10.8	17.69	*		-	-
15	PLACE4000466	13.01	11.54	9.57	6.56	6.92	4.59	5.76	4.65	5.6	*	**	-	-
	PLACE4000472	165.28	158.69	152.44	150.34	164.09	112.33	151.06	116.56	147.01			-	-
	PLACE4000487	10.38	6.99	9.12	4.68	4.63	2.93	3.52	2.06	3.49	*	**	-	-
	PLACE4000489	6.8	7.97	5.85	3.99	5.8	4.43	4.22	4.11	4.53	*		-	-
	PLACE4000494	70.51	55.91	43.58	29.17	31.84	24.64	72.67	73.79	78.08	*		-	-
20	PLACE4000502	204.58	180.8	176.1	156.92	166.08	118.92	216.32	165.78	200.53			-	-
	PLACE4000521	28.26	22.58	17.07	7.12	15.76	9	21.36	20.06	19.61	*		-	-
	PLACE4000522	18.46	20.38	16.27	10.46	14.51	9.51	16.91	13.88	17.23	*		-	-
	PLACE4000537	9.79	11.19	9.86	4.99	8.2	4.43	2.5	3.6	0.68	*	**	-	-
	PLACE4000548	12.35	10.52	9.27	3.9	4.07	4.41	4.08	3.04	3.64	**	**	-	-
	PLACE4000558	56.63	61.59	46.82	23.74	32.06	21.71	66.44	39.24	52.93	**		-	-
25	PLACE4000581	9.34	8.43	8.7	5.21	3.71	3.17	5.23	4.69	4.75	**	**	-	-
	PLACE4000590	4.28	4.39	2.54	2.29	1.9	1.58	2.69	1.47	1.72	*		-	-
	PLACE4000593	9.43	8.29	6.55	6.59	7.86	5.28	9.13	5.16	4.1			-	-
	PLACE4000612	50.54	44.64	46.86	32.73	32.12	18.75	8.58	0	9.68	*	**	-	-
	PLACE4000638	8.47	6.12	6.11	6.07	5.18	3.4	1.44	4.91	1.37	*		-	-
	PLACE4000650	11.71	6	6.58	1.78	5.39	2.69	5.01	3.89	2.3			-	-
30	PLACE4000651	10.02	10.01	7.86	4.63	8.67	3.43	7.21	6.04	5.23	*		-	-
	PLACE4000654	40.79	37.09	28.74	21.7	29.39	18.93	39.41	24.87	38.02			-	-
	PLACE4000670	7.33	7.03	6.77	5.23	9.34	3.97	6.41	3.63	6.87			-	-
	PLACE4000685	27.36	21.25	28.71	7.25	13.45	14.89	8.25	13.79	8.57	*	**	-	-
	PLACE4000687	5.87	2.72	5.94	3.75	3.52	3.04	2.8	3.66	2.1			-	-
	PLACE5000003	13.69	15.66	16.08	8.92	15.85	13.17	6.36	9.93	8.84	**		-	-
35	PLACE5000005	17.05	13.36	13.46	14.47	17.35	13.76	5.61	8.29	7.64	**	**	-	-
	PLACE5000019	9.12	11.38	11.73	6.9	5.65	5.55	3.51	3.39	6.14	**	**	-	-
	PLACE5000021	28.4	29.38	31.96	18.99	19.21	16.52	23.2	21.13	21.28	**	**	-	-
	PLACE5000022	44.41	69.47	62.06	68.78	41.05	40.85	49.9	41.64	44.86			-	-
	PLACE5000024	5.49	3.54	5.62	3.22	3.37	3.62	2.52	4.61	6.39			-	-
40	PLACE5000036	104.88	78.2	100.83	80.5	90.12	87.43	80.46	55.18	59.66			-	-
	PLACE5000059	13.88	12.58	10.61	5.28	9.26	6.17	6.41	11.67	8.18	*		-	-
	PLACE5000076	102.38	115.06	146.87	71.48	155.46	121.21	73.46	149.18	129.94			-	-
	PLACE5000117	10.83	17.74	16.15	14.79	20.06	15.63	5.73	9.53	5.57	*		-	-
	PLACE5000143	12.99	9.48	15.16	8.13	9.83	8.06	5.81	8.66	8.8			-	-
	PLACE5000152	183.88	202.8	96.28	147.85	104.4	116.95	74.92	70.73	75.92			-	-
	PLACE5000154	90.81	108.58	44.13	75.4	77.68	45.59	40.84	23.33	50.85			-	-
45	PLACE5000155	33.2	24.58	26.07	16.94	31.52	23.72	16.67	17.93	27.02			-	-
	PLACE5000165	113.75	76.52	82.55	90.32	103.03	88.36	32.72	55.94	39.36	*		-	-
	SKNMC1000004	1323.5	940.91	1083.3	685.71	1250.5	673.39	1094.9	1184.2	1060.5			-	-
	SKNMC1000011	8.28	9.06	11.81	4.48	12.47	5.45	4.79	5.44	8.03			-	-
	SKNMC1000013	14.72	15.01	20.62	10.84	11.72	12.41	4.74	6.54	6.23	**		-	-
50	SKNMC1000014	20.79	23.92	26.42	20.23	26.13	18.1	14.76	15.14	11.78	**		-	-
	SKNMC1000018	9.56	8.87	12.44	5.32	9.73	4.54	3.81	4.41	5.25	**		-	-
	SKNMC1000020	25.49	27.02	25.39	13.63	25.33	12.68	18.7	20.72	21.65	**		-	-
	SKNMC1000046	24.6	15.36	17.74	13.48	22.27	16.07	12.53	14.97	15.86			-	-
	SKNMC1000050	8.51	7.09	7.49	5.47	4.73	4.3	4.47	21.02	3.74	**		-	-
	SKNMC1000082	26.94	17.88	24.61	14.13	15.7	9.74	23.61	36.47	28.61	*		-	-
	SKNMC1000075	11.6	7.44	13.24	7.34	10.78	7.51	4.21	5.3	7.09			-	-
55	SKNMC1000082	22.01	16.02	26.11	8.38	13.04	9.62	12.8	21.17	16.8	*		-	-
	SKNMC1000091	36.26	39.39	43.7	12.98	23.22	18.68	24.3	27.26	22.42	**	**	-	-
	SKNMC1000098	11.34	8.86	15.64	6.38	7.5	4.4	5.83	7.15	6.38			-	-

Table 496

	PLACE3000400	42.78	33.52	30.29	11.43	26.51	8.47	52.76	33.47	38.46	*	*	-
	PLACE3000401	137.51	137.03	95.06	91.28	111.28	51.38	81.66	72.2	62.65	*	*	-
5	PLACE3000402	8.43	7.72	4.24	2.44	6.58	2.29	8.31	7.96	5.44			
	PLACE3000405	7.33	4.67	7.44	2.73	3.5	5.1	3.7	5.44	3.66			
	PLACE3000406	4.67	3.39	6.42	4.46	6.89	4.16	3.34	4.69	2.59			
	PLACE3000413	12.33	14.38	16.34	10.31	15.31	7.77	6.83	11.34	9.07	*	*	-
	PLACE3000416	84.6	58.17	41.71	94.16	126.08	86.52	51.61	89.65	63.01			
	PLACE3000425	2.87	4.19	4.03	2.67	4.11	2.12	2.08	0.77	1.54	*	*	-
10	PLACE3000437	6.08	4.62	9.74	3.57	11.5	3	3.57	3.67	4.77			
	PLACE3000455	6.66	7.45	7.23	5.48	5.38	4.19	5.5	5.74	3.82	*	*	-
	PLACE3000475	4.99	6.71	5.08	4.69	5.48	5.91	3.2	5.35	2.87			
	PLACE3000477	12.61	9.21	10.6	8.81	9.19	10.36	8.13	7.94	5.96	*	*	-
	PLACE4000003	30	20.47	39.94	8.43	18.43	9.82	12.11	10.52	8.23	*	*	-
	PLACE4000008	8.19	8.19	9.59	5.97	7.85	6.93	4.18	5.57	3.47	**	**	-
15	PLACE4000009	46.86	36	37.99	37.22	51.39	35.97	11.09	18.85	7.53	**	**	-
	PLACE4000014	18.34	18.57	19.39	16.62	19.49	16.59	6.08	13.2	15.87			
	PLACE4000029	5.3	5.07	5.95	2.91	4.87	2.62	3.69	3.02	2.87	**	**	-
	PLACE4000034	10.02	8.01	6.28	5.12	7.93	4.55	5.88	5.27	5.14			
	PLACE4000049	21.86	18.28	18.64	14.38	19.7	19.05	12.54	13.96	11.06	**	**	-
20	PLACE4000052	47.63	33.03	40.39	17.99	15.08	20.28	24.01	22.07	24.29	**	*	-
	PLACE4000062	53.46	46.6	50.02	32.69	40.48	28.08	39.71	36.99	45	*	*	-
	PLACE4000063	56.25	68.62	67.5	55.27	69.37	73.66	45.78	60.53	44.9			
	PLACE4000089	10.35	13.86	13.41	10.66	10.48	9.25	5.68	5.4	5.66	**	**	-
	PLACE4000093	15.33	18.02	19.57	10.49	13.84	10.33	17.17	14.6	14.04	*	*	-
	PLACE4000100	8.21	7.32	11.21	7.61	6.2	7.32	5.19	5.2	4.71	*	*	-
25	PLACE4000103	13.85	15.73	12.84	8.67	14.38	6.43	4.02	8.73	6.65	**	**	-
	PLACE4000106	32	18.81	23.23	18.35	21.38	15.86	17.78	20.58	18.06			
	PLACE4000128	22.05	18.53	23.14	8.88	10.24	9.57	11.18	17.49	11.76	**	*	-
	PLACE4000129	266.34	156.01	188.25	120.37	95.41	112.42	146.64	98.31	136.33	*	*	-
	PLACE4000131	59.1	45.51	65.57	41.68	34.82	39.78	18.29	19.7	30.83	*	**	-
	PLACE4000147	7.89	7.51	11.76	3.51	5.07	3.96	2.77	6.32	2.52	*	*	-
30	PLACE4000156	15.48	18.75	19.37	9.1	12.85	10.65	15.8	15.48	10.02	*	*	-
	PLACE4000175	12.45	10.41	18.08	11.94	9.67	2.46	6.38	6.69	6.89	*	*	-
	PLACE4000190	40.84	40.93	32.85	23.06	35.22	26.66	12.42	10.73	11	**	**	-
	PLACE4000192	35.69	24.13	23.88	16.77	21.59	17.55	17.75	19.51	19.4			
	PLACE4000206	33.82	29.03	26.48	11.73	17.97	14.5	17.45	20.16	20.02	**	*	-
	PLACE4000211	12.98	13.88	13.86	7.96	9.94	9.19	6.64	31.04	6.54	**	*	-
35	PLACE4000214	13.12	6.23	9.29	3.45	7.5	4.16	4.96	7.25	13.71			
	PLACE4000222	35.35	30.73	34.54	24.03	24.44	20.1	20.58	28.37	25.45	**	*	-
	PLACE4000223	14.88	14.83	15.03	7.36	10.28	5.52	4.16	7.67	5.85	**	**	-
	PLACE4000229	12.38	12.15	12.52	4.51	6.31	4.98	4.32	4.28	3.98	**	**	-
	PLACE4000230	9.56	8.77	9.56	4.68	7.06	5.7	4.87	9.82	3.71	**	*	-
	PLACE4000233	22.71	24.01	29.85	11.57	13.98	13.77	8.53	11.22	9.26	**	**	-
40	PLACE4000239	10.07	7.4	9.42	6.8	7.34	6.09	4.49	4.54	3.54	**	*	-
	PLACE4000247	15	11.82	15.56	9.41	13.91	8.87	8.67	6.24	7.39	**	*	-
	PLACE4000250	16.14	18.27	20.25	16.54	17.86	12.73	10.47	8.04	11.21	**	*	-
	PLACE4000252	10.01	5.15	12.75	1.39	3.96	3.1	2.21	3.92	2.7	*	*	-
	PLACE4000259	25.72	27.31	16.16	12.78	19.39	8.44	8.29	19.39	9.08			
	PLACE4000261	23.52	25.16	22.9	11.05	14.2	9.22	13.61	14.62	13.29	**	**	-
45	PLACE4000264	176.03	156.54	118.76	135.7	191.67	147.27	105.16	95.1	89.72	*	*	-
	PLACE4000269	71.18	49.07	62.08	23.03	41.63	24.74	71.71	82.6	52.18	*	*	-
	PLACE4000270	7.6	6.37	8.9	4.69	5.92	4.81	3.06	3.68	3.44	*	**	-
	PLACE4000281	15.26	16.07	18.29	6.09	10.92	7.61	6.44	6.66	5.1	**	**	-
	PLACE4000300	8.43	9.82	8.78	6.3	8.5	7.03	3.68	2.93	2.96	*	**	-
	PLACE4000320	12.37	11.22	11.92	6.27	10.09	6.43	4.47	4.26	2.85	*	**	-
50	PLACE4000323	15.16	16.25	19.16	9.12	14.06	10.9	6.05	4.54	4.03	*	**	-
	PLACE4000326	11.15	8.97	10.53	4.35	4.69	4.13	4.28	3.68	4.65	**	**	-
	PLACE4000344	27.71	32.93	24.51	14.27	30.44	19.64	5.14	9.78	8.98	**	*	-
	PLACE4000347	156.82	129.54	154.28	89.79	129.04	79.82	138.06	137.09	135.75			
	PLACE4000354	15.47	9.46	11.43	7.01	8.32	6.55	3.28	5.96	4.83	*	*	-
	PLACE4000367	8.63	5.47	7.82	5.38	5.39	6.72	3.84	3.95	3.64	*	*	-
55	PLACE4000369	11.27	6.37	9.31	5.27	4.5	4.05	4.11	4.32	3	*	*	-
	PLACE4000379	6.35	6	4.77	3.22	2.93	2.33	3.1	3.32	2.21	**	**	-
	PLACE4000387	25.99	34.88	32.06	17.28	20.68	16.67	6.75	8.86	3.97	*	**	-

Table 495

	PLACE3000067	49.68	45.52	57.37	38.78	41.13	26.16	39.8	46.08	41.69				
	PLACE3000069	22.6	20.15	23	9.32	14.75	10.81	1.9	9.3	14.16	**	*	-	-
	PLACE3000070	28.36	29.24	22.17	14.93	25.34	18.73	8.92	18.55	16.21		*	-	-
5	PLACE3000103	7.35	7.9	9.44	5.25	4.6	6.13	2.42	4.32	5.29	*	*	-	-
	PLACE3000119	35.89	28.1	30.76	15.16	14.79	15.09	26.05	30.71	30.91	**		-	-
	PLACE3000121	42.11	31.92	37.82	38.13	38.16	27.02	30.65	23.15	27.55			-	-
	PLACE3000124	419.8	380.05	314.65	379.74	412.05	282.09	411.56	374.59	344.25			-	-
	PLACE3000135	20.62	25.87	26.77	15.03	9.64	8.12	18.24	11.89	14.6	**	*	-	-
10	PLACE3000136	7.68	8.11	13.16	2.72	3.39	4.18	4.52	3.19	1.88	*	*	-	-
	PLACE3000142	24.15	28.81	26.5	11.04	24.59	15.08	9.37	6.94	11.56	**		-	-
	PLACE3000145	16.45	12.13	16.32	7.23	16.52	6.93	8.46	9.12	6.76	*		-	-
	PLACE3000147	109.42	62.46	138.9	40.47	87.42	32.89	57.71	63.5	68.14			-	-
	PLACE3000148	66.44	59.97	67.42	62.74	65.96	38.34	37.63	26.14	36.79	**		-	-
	PLACE3000154	10.3	11.14	12.88	7.95	8.64	6.56	7.31	6.3	7.57	*	**	-	-
15	PLACE3000155	54.28	50.77	22.77	23.49	28.34	24.7	10.12	11.93	8.34	*		-	-
	PLACE3000156	9.77	6.45	8.44	2.67	5.36	4.07	2.62	2.16	2.62	*	**	-	-
	PLACE3000157	38.44	50.52	36.49	23.64	29.36	25.04	21.36	24.91	20.01	*	*	-	-
	PLACE3000158	10.22	9.34	10.13	4.45	4.83	3.36	2.93	2.25	3.02	**		-	-
	PLACE3000160	19.92	29.65	15.93	8.71	11.7	10.32	5.77	7.29	4.57	*	*	-	-
	PLACE3000169	4.65	6.89	7.95	4.88	4.57	2.21	2.88	2.72	3.49	*		-	-
20	PLACE3000181	12.63	9.87	13.46	6.15	6.69	4.33	5.58	7.04	5.47	**	**	-	-
	PLACE3000194	1627	2636.2	1913.4	1710.4	2606.8	2399	3320.1	2826.5	2357.1			-	-
	PLACE3000197	7.18	4.55	8.38	2.14	1.71	1.66	2.79	2.24	2.57	*	*	-	-
	PLACE3000199	19.46	15.17	17.17	4.98	8.73	7.24	6.33	5.56	6.07	**	**	-	-
	PLACE3000205	12.53	16.93	16.84	6.91	6.83	4.67	7.28	5.87	4.15	**	**	-	-
	PLACE3000207	7.97	6.69	7.83	2.48	4.45	2.85	2.71	2.19	2.08	**	**	-	-
25	PLACE3000208	5.49	5.5	6.02	1.68	6.49	2.8	2.64	3.52	2.46	**		-	-
	PLACE3000213	40.02	27.48	31.62	27.29	30.91	23.89	26.59	30.58	23.2			-	-
	PLACE3000215	20.76	15.1	17.45	10.19	8.64	8.25	15.48	14.18	17.02	**		-	-
	PLACE3000218	13.39	10.52	10.44	6.03	3.81	4.29	4.65	4.57	4.68	**	**	-	-
	PLACE3000220	61.41	56.02	42.84	39.7	37	28.53	36.15	41.5	44.54	*		-	-
	PLACE3000221	12.37	9.42	8.54	5.7	3.63	2.33	5.09	3.54	2.87	*	**	-	-
30	PLACE3000225	180.89	144.07	131.66	176.14	175.29	127.3	77.77	64.65	86.2	**		-	-
	PLACE3000226	20.28	21.59	18.74	7.11	11.09	6.3	3.48	2.23	3.79	**	**	-	-
	PLACE3000230	17.46	18.98	14.94	6.27	14.11	10.52	4.65	3.25	5.88	**		-	-
	PLACE3000231	17.8	13.66	8.65	9.08	2.29	4.95	6.33	9.35	7.91			-	-
	PLACE3000235	8.61	7.55	8.28	4.48	5.65	4.02	5.72	2.93	4.77	**	*	-	-
	PLACE3000242	58.11	27.4	48.85	17.82	13.76	17.4	11.61	10.29	11.25	*	*	-	-
35	PLACE3000244	18.63	13.16	20	8.5	7.54	7.64	10.16	10.96	16.61	*		-	-
	PLACE3000253	56.99	47.97	42.85	32.37	25.93	24.8	32.98	24.76	32.96	*	*	-	-
	PLACE3000254	27.82	25.4	19.01	15.47	10.42	14.64	15.57	11.42	13.74	*	*	-	-
	PLACE3000271	14.58	20.8	13.28	5.71	8.52	5.71	8.32	6.43	8.38	*	*	-	-
	PLACE3000276	13.09	13.68	8.83	2.78	7.9	3.42	3.8	5.28	3.1	*	**	-	-
	PLACE3000304	37.31	49.14	39.5	28.72	17.9	18.26	33.17	28.5	40.23	*		-	-
40	PLACE3000308	12.29	9.64	9.42	6.2	5.6	4.27	5.42	4.79	6.47	**	**	-	-
	PLACE3000310	51.34	49.06	45.31	11.5	35.47	14.78	19.66	9.79	16.36	*	**	-	-
	PLACE3000320	24.3	28.51	23.33	14.31	20.51	12.27	15.7	15.03	15.45	*	*	-	-
	PLACE3000322	43.24	35.12	33.62	31.55	31.68	23.75	27.52	20.55	23.56	**		-	-
	PLACE3000330	28.37	27.71	24.49	12.33	15.5	11.35	20.66	17.37	21.33	**	*	-	-
	PLACE3000331	6.68	8.07	7.67	6.55	5.13	3.84	2.57	2.99	0.04	**		-	-
45	PLACE3000336	393.24	473.08	270.29	359.87	515.03	306.41	470.46	662.39	717.67			-	-
	PLACE3000338	11.78	5.98	6.82	3.79	5.41	2.48	8.74	10.29	11.53			-	-
	PLACE3000341	12.2	6.88	6.99	4.2	5.65	4.02	4.35	5.75	4.94			-	-
	PLACE3000350	13.71	12.66	10.21	4.86	8.13	4.84	5.77	4.37	7.49	*	*	-	-
	PLACE3000352	17.23	29.11	20.8	9.42	15.27	8.16	11.43	7.14	13.58	*		-	-
	PLACE3000353	14.74	14.96	16.82	8.78	8.26	6.98	10.37	5.48	6.79	**	**	-	-
50	PLACE3000362	57.42	93.8	72.53	67.02	59.59	35.38	59.68	38.27	60.6			-	-
	PLACE3000363	9.44	8.61	10.55	4.39	5.19	4.3	7.06	4.68	6.56	**	*	-	-
	PLACE3000365	5.59	7.45	4.94	1.48	2.65	3.2	4.78	2.37	2.95	*		-	-
	PLACE3000373	38.45	30.27	25.67	4.05	7.88	3.63	14.16	11.37	15.74	**	*	-	-
	PLACE3000374	48.65	44.25	34.93	16.2	26.01	11.78	24.23	27.25	23.51	*	*	-	-
55	PLACE3000387	5.55	3.77	3.38	1.58	3.57	0.43	3.58	2.95	3.21			-	-
	PLACE3000388	26.03	22.21	15.9	9.86	12.79	6.37	21.94	26.7	24.22	*		-	-
	PLACE3000398	15.37	13.07	9.83	5.64	4.9	3.93	10.68	11.05	9.52	**		-	-

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	PLACE2000172	9.55	12.02	10.86	4.88	5.87	3.32	4.01	4.8	6.2	**	**	-	-
	PLACE2000173	16.63	20.86	19.04	9.66	11.13	9.7	6.74	6.23	3.14	**	**	-	-
5	PLACE2000174	6.73	7.76	6.4	3.37	4.73	2.4	3.08	3.26	1.62	*	**	-	-
	PLACE2000176	47.07	39.92	36.01	26.8	32.78	24.97	24.68	26.79	25.41	*	**	-	-
	PLACE2000187	11.49	9.44	7.22	4.28	4.58	4.45	2.47	4.64	3.94	*	*	-	-
	PLACE2000216	15.24	13.28	16.1	7.83	10.53	7.67	11.12	11.23	10.67	**	*	-	-
	PLACE2000219	43.32	28.21	37.3	18.17	6.67	13.94	7.68	7.79	6.4	*	**	-	-
	PLACE2000221	9.58	7.64	7.46	2.22	3.99	1.24	4.08	2.43	3.07	**	**	-	-
10	PLACE2000223	16.67	12.34	12.95	4.95	6.75	4.32	4.43	2.01	2.87	**	**	-	-
	PLACE2000231	8.48	8.65	7.01	3.78	5.92	4.27	4.19	2.32	3.59	*	**	-	-
	PLACE2000235	7.02	5.03	3.66	1.54	5.33	1.9	4.43	3.57	3.78			-	-
	PLACE2000246	21.31	17.8	13.21	7.9	13.4	6.47	11.52	9.6	10.91	*		-	-
	PLACE2000264	41.11	43.3	30.68	20.28	22.27	18.14	15.54	15.85	15.69	*	**	-	-
	PLACE2000274	6.8	7.42	7.36	4.35	3.04	1.89	3.84	3.12	4.31	**	**	-	-
15	PLACE2000287	32.11	30.76	24.99	20.39	25.61	21.53	17.96	15.3	19.45	**		-	-
	PLACE2000296	42.94	34.88	35.63	23.36	22.17	20.37	32.02	14.36	29.35	**		-	-
	PLACE2000302	252.74	304.77	237.45	265.35	267.99	217.29	124.09	112.58	108.48	**	**	-	-
	PLACE2000305	21.47	23.58	17.69	16.14	17.23	16.65	22.14	18.21	22.01			-	-
	PLACE2000317	37.2	38.12	23.24	20.01	28.38	14.37	31.05	33.16	31.64			-	-
20	PLACE2000324	18.44	17.84	10.35	6.13	8.75	4.98	3.24	8.74	3.2	*	*	-	-
	PLACE2000334	21.38	17.6	11.43	10.36	9.7	9.37	7.54	6.15	9.1	*	*	-	-
	PLACE2000335	71.59	66.59	54.82	51.15	35.99	40.86	33.77	36.69	29.67	*	**	-	-
	PLACE2000340	20.62	28.95	19	17.88	23.88	12.85	16.23	14.73	16.82			-	-
	PLACE2000341	14.8	11.92	10.45	5.66	5.34	5.33	5.38	5.46	7.66	**	*	-	-
	PLACE2000342	23.53	21.95	21.97	16.28	11.86	11.51	22.81	14.69	23.12	**		-	-
25	PLACE2000347	9.57	9.26	10.05	4.16	2.94	3.3	5.78	3.54	5.48	**	**	-	-
	PLACE2000357	112.17	70.59	69.73	83.45	100.62	78.25	23	68.79	48.54			-	-
	PLACE2000358	57.37	32.59	34.74	25.22	20.69	15.46	44.93	47.18	42.01			-	-
	PLACE2000359	66.39	48.68	57.2	40.91	54.54	32.45	1.96	27.1	55			-	-
	PLACE2000366	24.09	23.36	19.56	12.65	12.71	9.05	11.75	9.72	10.83	**	**	-	-
	PLACE2000371	12.17	9.37	8.13	5.98	7.76	4.49	7.64	10.97	6.2			-	-
	PLACE2000373	8.3	7.58	8.83	4.52	6.25	4.81	3.89	1.17	2.44	**	**	-	-
30	PLACE2000374	20.82	21.57	18.21	16.3	16.71	14.02	11.71	11.44	14.19	*	**	-	-
	PLACE2000379	109.97	110.55	125.02	99.68	141.5	89.14	84.64	65.75	71.64	**	**	-	-
	PLACE2000386	27.43	27.19	20.99	10.83	12.4	11.22	15.3	17.99	10.26	**	*	-	-
	PLACE2000388	250.97	170.84	203.4	172.85	236.63	173.11	130.31	109.84	93.57	*		-	-
	PLACE2000392	241.74	206.31	271.34	159.33	199.67	145.83	119.57	115.9	95.8	*	**	-	-
35	PLACE2000394	10.9	10.42	13.61	8.59	15.25	7.68	5.68	9.12	6.51	*	*	-	-
	PLACE2000398	16.83	19.82	17.04	9.21	12.98	10.99	7.5	15.14	8.57	**	*	-	-
	PLACE2000399	30.6	27.17	17.03	25.83	27.41	28.06	9.08	19.74	20.17			-	-
	PLACE2000402	5.25	8.04	8.31	4.2	9.81	3.39	4.84	3.89	6.41			-	-
	PLACE2000404	24.79	49.52	29.14	49.74	21.66	25.22	16.23	19.06	8.51			-	-
	PLACE2000411	5.67	8.96	6.76	7.21	11.84	6.13	5.19	4.54	7.08			-	-
40	PLACE2000418	10.3	8.94	9.67	6.1	3.09	4.67	4.68	5.3	4.11	**	**	-	-
	PLACE2000419	96.71	94.48	148.23	58.48	57.97	27.06	44.37	46.85	45.47	*	*	-	-
	PLACE2000425	9.83	11.3	13.49	7.49	11.26	6.24	5.84	7.49	5.62	*		-	-
	PLACE2000427	15.21	22.92	21.67	15.7	21.56	15.92	5.61	6.9	4.76	**		-	-
	PLACE2000433	25.58	27.32	26.12	13.97	18.4	14.15	21.62	27.64	18.08	**		-	-
	PLACE2000435	16.36	18.11	21.19	9.69	13.76	7.6	8.43	10.79	12.73	*	*	-	-
45	PLACE2000438	35.17	36.25	36.72	24.14	34.38	15.98	38.19	37.23	25.32			-	-
	PLACE2000450	24.38	41.59	23.11	16.35	37.33	18.21	23.47	16.65	30.25			-	-
	PLACE2000455	51.05	47.04	53.6	30.31	36.57	21.71	39.12	34.02	35.42	*	**	-	-
	PLACE2000458	13.07	11.58	13.21	6.82	7.75	7.03	6.6	8.78	5.76	**	**	-	-
	PLACE2000464	9.95	10.55	11.94	7.59	6.37	5.51	7.66	8.47	7.16	**	*	-	-
	PLACE2000465	182.2	189.36	208.18	166.02	148.04	156.91	139.29	234.97	151.61	*		-	-
50	PLACE2000473	23.66	24.51	24.43	18.65	18.15	16.44	24.18	30.07	29.1	**		-	-
	PLACE2000477	14.25	12.47	18.55	9.69	12.47	6.23	10.6	9.09	9.9	*		-	-
	PLACE3000004	48.49	41.47	56.9	42.63	53.35	39.93	39.48	28.43	32.45	*		-	-
	PLACE3000009	30.13	36.74	28.18	19.79	35.05	20.31	13.54	15.44	20.29	**		-	-
	PLACE3000020	27.99	24.44	23.74	13.45	15.64	8.65	14.29	14.44	13.63	**	**	-	-
	PLACE3000029	13.95	12.7	16.35	6.3	9.27	7.3	4.68	16.59	5.52	**		-	-
55	PLACE3000038	6.52	4.32	7.69	5.63	3.97	4.12	2.68	6.95	6.18			-	-
	PLACE3000052	8.21	5.21	10.48	3.53	4.29	4.15	2.75	5.07	2.29			-	-
	PLACE3000058	7.39	8.1	10.8	4	5.08	2.55	2.83	4.28	4.31	*	*	-	-

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	PLACE1011836	18.92	14.13	10.67	5.56	9.72	5.44	13.34	9.9	14.01				
	PLACE1011847	18.99	16.08	12.62	9.39	11.06	5.84	18.15	10.26	13.1	*		-	
5	PLACE1011855	44.57	41.99	25.3	17.11	33.33	14.76	55.6	42.68	40.57				
	PLACE1011858	17.26	16.38	17.11	5.33	7.08	6.53	6.22	8.74	5.1	**	**	-	-
	PLACE1011874	23.12	12.72	25.74	16.84	19.22	7.85	10.17	8.71	12.51				
	PLACE1011875	10.58	9.98	11.17	5.18	10.63	6.87	7.18	6.04	4.93		**	-	
	PLACE1011877	4.39	5.14	4.95	3.7	5.52	3.78	2.59	4.05	4.24				
	PLACE1011891	21.71	17.57	18.99	7.39	14.47	8.67	8.21	9.15	7.86	*	**	-	-
10	PLACE1011895	15.35	16.64	24.3	10.41	10.14	12.14	5.01	8.13	7.98	*		-	-
	PLACE1011920	16.12	14.42	15.9	7.68	8.85	7.76	6.46	2.46	3.24	**	**	-	-
	PLACE1011922	9.14	7.55	8.03	4.49	8.39	4.08	7.97	6.15	5.33				
	PLACE1011923	166.86	139.43	146.9	104.2	117.42	86.95	136.93	140	124.61	*		-	-
	PLACE1011937	11.45	9.18	10.8	8.58	6.75	5.1	5.71	6.29	5.46	*	**	-	-
	PLACE1011939	20.33	21.55	26.93	12.79	15.19	13.44	14.58	11.31	12.67	*	*	-	-
15	PLACE1011940	46.38	48.76	46.73	35.11	34.53	31.67	43.31	41.44	39.86	**	**	-	-
	PLACE1011962	58.59	70.73	80.18	30.15	38.09	28.84	47.96	47.68	42.57	**	*	-	-
	PLACE1011964	53.78	49.36	46.95	41.43	47.33	36.34	22.23	20.62	20.74	**		-	-
	PLACE1011978	50.21	52.86	64.71	38.6	31.39	22.36	22.46	20.19	18.12	*	**	-	-
	PLACE1011980	13.57	18.54	10.96	9.65	15.28	6.53	4.68	9.99	8.31				
	PLACE1011981	46.23	28.89	45.44	16.98	15.61	9.98	14.66	11.01	11.18	*	**	-	-
20	PLACE1011982	116.69	99.4	123.56	57.42	68.6	45.01	97.45	90.55	115.07	**		-	-
	PLACE1011995	8.31	11.27	12.33	5	7.29	7.22	6.27	8.29	4.87	*		-	-
	PLACE1012023	36.36	36.13	39.16	21.46	26.03	23.3	20.6	21.06	24.79	**	**	-	-
	PLACE1012026	72.96	80.46	66.75	52.71	40.29	50.42	21.72	33.87	25.71	**	**	-	-
	PLACE1012031	6.69	7.34	9.86	2.73	6.11	3.89	3.41	3.11	3.97	*		-	-
25	PLACE2000003	144.75	222.75	62.64	74.47	133.88	40.79	39.01	32.21	30.64				
	PLACE2000005	20.72	25.78	18.16	9.79	22.43	11.52	5.79	8.65	14.3	*		-	-
	PLACE2000006	74.92	51.78	61.3	27.4	24.75	21.12	50.02	43.94	72.77	**	*	-	-
	PLACE2000007	11.31	11.29	12.53	6.68	7.02	5.2	5.63	10.02	7	**	*	-	-
	PLACE2000011	20.07	19.36	23.84	14.92	4.58	15.77	9.39	11.52	19.43				
	PLACE2000014	10.47	6.72	12.3	4.76	6.02	3.48	3.63	6.9	4.73	*		-	-
30	PLACE2000015	63.77	83.22	43.62	34.06	43.14	43.39	19.29	26.64	36.8	*		-	-
	PLACE2000017	64.44	66.07	78.07	40.68	48.09	29.95	59.7	61.53	44.91	*		-	-
	PLACE2000021	131	136.36	142.32	116.28	142.66	101.82	89.75	68.71	86.35	**	**	-	-
	PLACE2000022	6.52	6.75	7.52	3.85	5.14	3.55	4.53	1.84	5.72	**		-	-
	PLACE2000030	277.16	275.48	252.39	182.81	167.16	156.31	125.23	96.07	135.65	**	**	-	-
	PLACE2000032	8.03	6.4	8.8	4.38	5.33	2.81	4.32	15.18	4.01	*		-	-
35	PLACE2000033	19.24	16.61	20.22	9.43	14.03	10.83	8.93	12.91	10.87	*	**	-	-
	PLACE2000034	212.75	182.06	146.24	219.9	195.88	170.13	146.86	112.34	125.26				
	PLACE2000039	8.9	10.84	13.51	4.94	3.89	4.8	3.72	4.38	6.52	**	*	-	-
	PLACE2000043	13.64	20.44	15.11	6.28	13.12	6.04	6.05	7.64	6.22	*		-	-
	PLACE2000044	7.28	9.48	9.44	3.48	5.71	4.79	4.48	4.25	4.59	*	**	-	-
	PLACE2000047	17.41	19.39	12.83	6.83	12.83	7.05	7.81	8.54	5.78	*		-	-
40	PLACE2000050	18.34	17.87	22.05	7.34	8.98	7.31	7.51	8.31	7.94	**	**	-	-
	PLACE2000061	53.75	61.97	78.23	34.92	41.03	37.86	22.68	14.04	20.77	*	**	-	-
	PLACE2000062	44.19	44.8	62.08	41.54	45.17	33.58	38.88	38.06	40.24				
	PLACE2000072	216.69	181.79	181.98	200.73	185.27	186.23	111.76	113.44	105.53		**	-	-
	PLACE2000073	8.16	10.75	9.27	3.77	4.89	3.62	4.56	4.79	3.59	**	**	-	-
	PLACE2000097	120.94	128.07	120.9	148.48	131.74	105.73	90.14	87.25	86.79	**		-	-
45	PLACE2000100	13.48	14.19	15.99	6.02	8.77	5.56	4.8	3.51	6.24	**	**	-	-
	PLACE2000103	355.83	328.95	197.42	251.22	436.52	235.01	231.23	336.03	237.95				
	PLACE2000106	32.23	23.41	23.14	14.46	6.17	13.03	9.13	10.5	5.27	*	**	-	-
	PLACE2000111	152.37	157.38	125.9	88.6	105.91	83.66	90.52	44.23	69.19	*	**	-	-
	PLACE2000115	27.01	16.97	16.32	19.6	21.67	16.26	13.74	11.32	11.63				
	PLACE2000118	229.23	214.41	143.27	124.01	152.19	92.15	153.7	62.85	67.07				
50	PLACE2000124	7.33	5.71	5.57	2.12	1.78	1.55	2.01	1.56	1.09	**	**	-	-
	PLACE2000132	18.07	19.01	19.11	15.64	11.16	12.49	9.7	6.52	8.48	*	**	-	-
	PLACE2000136	33.6	40.68	26.33	13.95	42.68	22.54	9.98	17.8	18.28	*		-	-
	PLACE2000137	6.87	5.34	6.91	1.61	2.95	2.09	1.24	1.8	2.36	**	**	-	-
	PLACE2000140	11.37	11.51	12.2	8.07	9.45	9.49	11.48	11.25	10.8	**		-	-
	PLACE2000147	10.22	7.24	9.46	6.01	5.85	4.51	7.52	7.6	9.27	*		-	-
55	PLACE2000153	17.95	21.42	13.85	5.51	9.57	6.43	4.84	5.27	2.27	*	**	-	-
	PLACE2000164	8.53	12.11	9.44	3.18	4.26	4.12	4.98	3.97	2.13	**	**	-	-
	PLACE2000170	38.84	46.93	33	28	28.56	22.27	30.03	23.36	30.82	*		-	-

Table 492

	PLACE1011325	67.49	68.67	34.36	32.79	55.55	25.52	26.06	16.66	23.25	*	-
	PLACE1011332	37.6	32.99	20.69	24.89	22.84	17.06	26.8	25.99	39.36		
5	PLACE1011340	6.86	4.22	8.9	7.42	3.33	10.93	4.69	3.22	3.76		
	PLACE1011353	13.06	9.45	9.31	4.05	5.07	3.21	9.83	8.83	10.17	**	-
	PLACE1011360	7.09	6.29	7.09	-5.01	4.64	2.08	4.25	9.92	6.13	*	-
	PLACE1011364	36.2	48	46.69	18.82	32.17	16.48	29.76	23.08	23.29	*	*
	PLACE1011365	56.47	40.06	43.45	30.4	25.78	22.53	35.29	36.44	44.51	*	-
	PLACE1011371	8.03	11.85	15.3	3.37	4.12	3.15	3.67	7.88	4.18	*	-
10	PLACE1011375	7.04	6.27	10.93	2.06	3.1	2.3	1.72	3.77	5.21	*	-
	PLACE1011386	164.83	196.07	182.53	143.65	164.65	123.82	185.62	179.87	160.52		
	PLACE1011399	13.53	14.93	12.77	5.31	8.23	5.03	14.49	12.53	13.08	**	-
	PLACE1011406	24.35	18.36	25.14	17.24	17.93	13.01	14.05	16.63	13.24	*	-
	PLACE1011407	11.96	8.15	11.8	4.39	4.52	4.88	4.7	7.06	4.99	**	*
	PLACE1011419	45.6	33.74	47.01	28.72	27.55	25.25	33.12	24.14	29.66	*	-
15	PLACE1011433	12.4	6.05	13.11	5.89	9.23	4.52	4.03	6.39	4.28		
	PLACE1011440	22.91	21.43	16.68	17.18	6.65	6.34	4.22	2.79	3.13	**	-
	PLACE1011452	21.99	27.18	21.9	7.47	5.58	8.33	1.98	2.55	3.76	**	**
	PLACE1011465	8.59	10.4	8.63	4.43	3.74	3.61	3.27	3.42	2.9	**	**
	PLACE1011472	12.35	11.03	8.39	5.76	5.63	4.74	4.12	4.06	4.53	*	**
	PLACE1011477	6.89	4.15	8.42	3.82	3.1	2.49	5.68	5.36	7.03		
20	PLACE1011478	147.8	127.45	151.17	98.43	129.14	92.91	97.94	88.28	67.65	**	-
	PLACE1011492	8.86	4.94	8.71	5.63	3.6	3.44	4.78	2.62	3.91		
	PLACE1011498	11.86	13.27	11.87	6.42	7.11	5.44	7.49	6.76	6.34	**	**
	PLACE1011501	5.51	5	4.76	1.97	2.88	1.92	1.53	1.09	0.63	**	**
	PLACE1011503	8.36	7.18	8.72	3.16	3.42	2.41	3.54	1.96	1.63	**	**
	PLACE1011509	6.79	8.76	6.6	2.67	3.22	3.57	3.54	1.66	2.46	**	**
25	PLACE1011514	39.36	42.6	35.59	29.05	24.04	21.59	28.05	21.32	28.44	**	*
	PLACE1011516	2.25	2.2	3.02	1.65	1.39	0.92	2.22	1.19	1.28	*	-
	PLACE1011520	56.28	54.22	44.97	60.14	59.13	46.66	37.37	33.63	31.08	*	-
	PLACE1011538	3.92	2.67	3.99	1.41	2.25	0.89	1.63	0.52	1.52	*	*
	PLACE1011555	5.04	5.36	4.18	1.71	1.41	2.12	2.42	1.56	2.13	**	**
	PLACE1011561	40.83	32.97	31.05	28.42	20.97	80.97	15.06	18.06	18.45	**	-
30	PLACE1011563	10.59	8.43	6.69	4.39	3.31	1.28	2.74	2.46	2.05	*	**
	PLACE1011567	8.99	5.72	8.85	4.85	2.48	0.94	3.39	1.71	1.08	*	*
	PLACE1011569	17.38	15.57	12.39	8.98	14.17	5.41	4.61	3.24	6.59	**	-
	PLACE1011576	9.4	4.88	4.15	2.79	3.81	3.52	4.89	6.65	7.51		
	PLACE1011586	11.29	10.02	9.62	6.88	3.58	5.49	4.18	5.75	5.57	**	**
35	PLACE1011635	30.93	17.84	23.55	13.98	14.55	6.78	13.1	10.06	12.62	*	-
	PLACE1011641	17.21	13.31	15.5	11.83	10.3	8.13	12.16	8.31	8.02	*	*
	PLACE1011642	14.04	9.07	8.81	4.73	3.8	2.88	2.08	3.09	1.68	*	**
	PLACE1011643	21.88	15.91	14.62	9.69	7.73	4.5	12.37	9.92	7.66	*	*
	PLACE1011646	50.88	55.98	42.82	29.54	24.58	20.05	12.83	13.88	9.31	**	**
	PLACE1011649	101.51	209.33	140.08	111.98	127.38	61.19	33.86	36.5	31.67	*	-
40	PLACE1011650	21.23	16.3	17.27	6.34	7.58	5.35	13.14	9.77	13.14	**	*
	PLACE1011661	36.54	29.6	21.98	17.54	11.85	16.02	29.9	27.07	30.09	*	-
	PLACE1011664	4693.7	3704.8	3103.3	1925.9	2495.7	2690.1	3091.7	3779.5	2474.5	*	-
	PLACE1011672	14.58	19.19	17.05	4.82	11.54	6.52	8.71	6.37	8.66	*	**
	PLACE1011675	27.27	28.5	23.64	11.76	7.61	13.02	10.47	6.83	11.83	**	**
	PLACE1011682	31.95	30.83	21.79	24.47	12.69	9.16	19.26	16.69	17.56	*	-
45	PLACE1011708	240.54	265.47	223.19	186.79	250.35	199.96	147.11	130.97	144.28	**	-
	PLACE1011719	2115.1	1324.5	1826.8	1953.8	1265.9	1000.2	1428.6	860.26	1269.9		
	PLACE1011725	14.63	11.44	8.67	4.74	6.3	2.04	9.22	9.94	8.53	*	-
	PLACE1011729	17.92	12.45	10.92	5.48	6.85	3.57	9.59	6.92	10.12	*	-
	PLACE1011741	19.11	16.45	8.97	6.48	6.39	4.13	6.03	4.62	6.01	*	*
	PLACE1011749	198.63	245.38	188.1	98.37	139.73	93.5	146.64	106.95	169.56	*	-
50	PLACE1011757	55.97	64.31	51.13	22.6	27.43	20.99	59.9	47.96	65.48	**	-
	PLACE1011762	112.55	118.61	84.99	66.65	83.7	57.21	113.37	92.71	109.2	*	-
	PLACE1011778	14.17	14.57	14.58	9.09	8.23	7.54	9.58	8.34	17.29	**	-
	PLACE1011783	34.3	38.67	23.05	14.78	12.16	17.09	15.47	21.92	20.45	*	-
	PLACE1011795	26.86	17.86	14.49	8.41	7.89	4.41	13.16	9.37	14.56	*	-
	PLACE1011810	6.46	2.52	3.35	2.53	2.07	0.8	3.19	3.06	3.38		
55	PLACE1011824	86.62	55.93	50.79	29.92	21.66	12.01	21.62	24.56	19.64	*	*
	PLACE1011825	47.11	53.31	45.03	28.52	34.47	24.16	61.42	44.87	47	**	-
	PLACE1011835	12.82	11.42	11.32	5.55	7.84	4.84	7.39	9.98	8.67	**	*

Table 491

	PLACE1010882	17.06	13.54	8.42	6.17	5.69	5.82	7.13	5.03	5.37	*	-
	PLACE1010891	27.43	30.32	21.15	13.46	14.41	7.46	15.38	15.07	24.71	*	-
5	PLACE1010896	13.62	18.23	16.16	6.92	8.03	5.78	3.54	4.62	4.06	**	**
	PLACE1010900	66.6	75.65	67.62	55.68	59.03	45.47	67.12	53.92	39.14	*	-
	PLACE1010916	104.52	71.53	76.5	35.13	35.16	33.1	29.84	23.73	27.17	**	**
	PLACE1010917	14.21	14.67	11.27	7.82	5.59	7.41	7.69	7.06	8.76	**	**
	PLACE1010924	5.86	2.7	4.5	2.89	4.47	3.08	2.47	3.1	1.31		
	PLACE1010925	4.94	4.55	4.99	1.88	2.53	1.67	3.09	1.69	1.78	**	**
10	PLACE1010926	13.94	11.16	11.63	5.87	7.64	6.22	5.6	4.23	4.29	**	**
	PLACE1010942	207.77	196.21	175.46	147.15	130.24	102.36	197.42	120.16	123.73	*	-
	PLACE1010943	31.5	26.2	22.98	12.49	22.16	9	15.12	16.22	13.62	*	-
	PLACE1010944	83.2	92.75	68.43	51.82	78.63	47.87	87	58.47	73.16		
	PLACE1010947	33.76	22.26	22.3	12.84	13.89	8.22	22.66	11.99	13.89	*	-
	PLACE1010954	10.33	11.3	11.05	5.3	5.81	5.51	5.34	5.12	4.22	**	**
15	PLACE1010960	15.62	30.44	20.31	7.94	8.28	7.04	11.37	9.05	8.5	*	*
	PLACE1010965	13.17	14.34	9.79	7.54	4.78	3.78	4.15	4.73	5.18	*	**
	PLACE1010968	18.41	14.95	14.27	6.84	9.6	5.59	12.62	6.34	9.37	**	*
	PLACE1010978	57.65	54.74	36.05	40.68	34.09	13.7	16.85	10.36	20.32	*	-
	PLACE1010982	21.44	26.06	15.9	7.42	6.54	11.09	10.66	7.85	9.32	*	*
	PLACE1010990	327.98	344.55	256.15	258.07	367.78	240.66	208.45	171.08	173.44	*	-
20	PLACE1011017	100.16	59.4	54.99	23.72	27.4	15.8	91.94	75.46	99.74	*	-
	PLACE1011019	13.37	8.64	6.46	8.37	6.37	4.79	6.61	7.41	7.81		
	PLACE1011026	17.67	16.91	13.69	8.74	11.48	8.93	15.71	13.37	15.94	*	-
	PLACE1011032	5.5	8.88	7.51	2.84	5.75	2.99	4.5	3.38	3.21	*	-
	PLACE1011041	11.32	11.72	11.41	4.79	6.17	4.9	2.74	0	1.85	**	**
	PLACE1011046	30.68	27.07	24.09	20.73	22.77	14.24	25.88	22.96	30.06	**	**
25	PLACE1011054	10.75	10.03	9.15	3.68	6.42	4.38	4.21	3.5	2.56	**	**
	PLACE1011056	23.11	26.67	18.54	13.24	17.45	11.68	27.3	21.77	27.93	*	-
	PLACE1011057	5.67	3.31	3.16	3.73	4.11	8.46	6.33	3.05	6.91		
	PLACE1011059	10.04	10.53	7.99	8.15	6.14	3.72	11.95	6.55	7.6		
	PLACE1011066	19.4	17.96	11.33	7.08	9.45	9.52	6.53	6.86	9.19	*	*
30	PLACE1011087	9.28	7.31	5.97	5.75	7.03	4.1	2.13	5.18	0.49	*	-
	PLACE1011090	26.84	40.15	33.04	57.32	27.48	16.29	127.9	19.42	37.4		
	PLACE1011099	10.52	7.52	6.48	3.01	5.18	3.1	6.04	4.61	6.16	*	-
	PLACE1011109	43.09	16.58	13.76	6.5	19.94	4.84	12.2	7.21	13.97		
	PLACE1011114	17.98	24.04	19.89	13.12	19.15	8.13	12.09	13.25	14.96	*	-
	PLACE1011116	30.03	28.53	37.37	10.98	8.87	7	8.26	10.18	5.99	**	**
35	PLACE1011122	5.9	6.6	5.93	4.16	4.82	3.11	3.97	4.69	2.89	*	*
	PLACE1011133	6.47	4.35	6.51	3.81	3.93	3.81	2.72	5.81	2.91		
	PLACE1011134	5.48	5.99	6.34	4.28	4.27	3.64	3.72	3.86	3.76	**	**
	PLACE1011143	4.76	9.67	5.59	2.21	5.45	3.27	2.93	3.83	4.13		
	PLACE1011146	10.48	13.56	12.48	9.3	11.11	4.11	4.38	6.69	4.48	**	-
	PLACE1011160	7.71	9.99	6.71	4.98	7.77	3.25	6.45	4.47	3.78		
40	PLACE1011165	21.89	27.95	32.58	8.87	19.75	11.93	12.6	15.59	9.42	*	*
	PLACE1011181	5.25	3.37	6	2.06	4.49	2.29	2.61	4.09	3.85		
	PLACE1011185	60.3	36.78	36.17	37.68	34.15	24.31	17.32	33.87	29.05		
	PLACE1011186	25.07	30.45	27.42	12.69	20.01	15.83	19.93	27.9	21.4	*	-
	PLACE1011203	10.27	18.2	13.8	9.36	12.5	6.63	8.02	13.49	10.02		
	PLACE1011214	7.39	9.12	10.82	5.57	5.78	3.95	1.49	2.02	3.84	*	**
45	PLACE1011219	7.03	5.86	7.27	8.39	4.52	4.58	9.07	4.64	3.23		
	PLACE1011221	10.63	11.42	9.42	5.44	10.51	4.92	4.92	5.52	6.28	**	-
	PLACE1011229	13.76	13.72	12.78	9.23	7.48	4.86	6.43	7.1	4.99	**	**
	PLACE1011231	34.58	28.7	50.02	23.03	24.8	13.83	19.12	24.97	26.97		
	PLACE1011236	13.02	10.94	13.96	5.99	7.59	5.06	4.42	6.98	4.82	**	**
	PLACE1011247	571.8	409.73	639.71	528.63	738.01	566.23	553.23	542.58	267.58		
50	PLACE1011263	4.49	7.26	9.65	4.09	5.29	4.39	2.77	4.83	3.37		
	PLACE1011273	9.77	15.11	15.29	8	7.81	4.87	6.04	5.28	3.97	*	*
	PLACE1011276	21.08	26.71	20.88	9.67	19.17	11.18	9.31	5.1	7.74	**	-
	PLACE1011289	8.76	9.56	9.49	5.47	7.47	4.38	4.05	4.14	4.61	*	**
	PLACE1011291	3.27	1.73	2.37	1.38	1.37	0.09	2.52	4.22	1.75		
	PLACE1011296	15.1	14.79	20.48	7.86	6.09	7.28	12.72	18.07	15.96	**	-
55	PLACE1011310	49.57	40.91	47.27	26.08	24.65	29.49	26.25	31.46	34.3	**	*
	PLACE1011311	12.85	11.18	13.64	5.84	6.4	7.88	4.02	7.29	3.99	**	**
	PLACE1011321	316.11	225.18	251.29	230.23	241.81	208.08	340.56	284.43	231.07		

Table 490

	PLACE1010364	11.29	7.43	7.22	2.26	3.51	2.16	4.95	3.65	4.16	*	*	-	-
	PLACE1010368	10.78	8.61	7.03	3.98	6.56	5.19	7.58	7.15	5.73				
5	PLACE1010373	30.2	28.61	24.17	14.82	22.58	10.46	24.63	20.09	20.87	*		-	
	PLACE1010383	18.56	18.15	16.04	10	15.54	3.52	27.13	12.41	12.03				
	PLACE1010385	2.48	1.73	1.18	0.83	2.33	0.49	2.65	1.19	2.06				
	PLACE1010389	8.48	7.51	8.42	3.76	4.47	5.47	3.96	4.51	2.91	**	**	-	-
	PLACE1010401	3.24	0.96	4.6	2.29	2.49	1.63	2.81	2.31	1.6				
	PLACE1010410	22.81	15.39	21.28	8.31	20.95	12.79	12.79	10.9	11.3	*	*	-	-
10	PLACE1010418	18.78	19.01	17.17	5.75	13.45	6.67	11.25	18.37	11.11	*		-	
	PLACE1010425	10.58	10.75	18.36	4.2	4.88	3.19	8.87	11.17	8.75	*		-	
	PLACE1010443	12.48	14.66	16.77	5.6	14.99	6.09	8.79	8.23	11.76	*		-	
	PLACE1010445	36.17	45.7	37.43	17.56	20.28	18.29	39.23	32.46	41.56	**	*	-	-
	PLACE1010481	60.4	46.95	65.37	22.92	26.42	21.44	27.71	30.53	27.4	**	**	-	-
	PLACE1010482	127.4	82.31	84.53	104.64	120.51	109.85	62.48	50.87	49.97	*		-	
15	PLACE1010491	71.28	72.3	73.41	43.51	47.95	35.09	68.63	86.51	74.84	**	*	-	-
	PLACE1010492	20.32	20.74	19.72	20.82	27.04	23.27	9.07	12.55	10.15	**	**	-	-
	PLACE1010509	11.2	13.07	17.83	8.18	12.58	8.24	5.79	8.34	7.68	*	*	-	-
	PLACE1010518	9.76	12.31	18.67	8.17	8.57	7.5	8.04	6.33	6.73				
	PLACE1010522	8.32	9.35	9.63	6.71	11.42	6.63	6.37	5.88	7.65	*		-	
	PLACE1010529	13.09	22.15	14.64	11.31	17.05	8.58	7.61	9.45	9.29				
20	PLACE1010547	36.79	34.33	38.34	12.39	16.81	11.19	21.21	24.3	23.05	**	**	-	-
	PLACE1010560	10.15	9.34	9.56	6.09	4.54	5.89	6.02	5.35	6.13	**	**	-	-
	PLACE1010562	4.79	4.39	12.01	8.8	8.65	6.38	6.41	7.48	4.4				
	PLACE1010579	74.54	67.98	59.08	46.33	48.92	48.24	56.27	84.49	60.31	*		-	
	PLACE1010580	38.79	45.7	46.75	22.24	30	17.17	31.95	38.27	36.2	*		-	
	PLACE1010599	29.35	25.54	32.71	12.61	15.12	14.18	19.83	24.34	23.16	**	*	-	-
25	PLACE1010606	31.76	30.37	40.62	15.46	19.15	14.23	22.77	25.83	28.81	**	*	-	-
	PLACE1010616	16.39	26.11	18.04	12.86	17.78	7.33	3.49	3.67	4.92	**	*	-	-
	PLACE1010622	9.08	4.75	12.08	2.52	4.26	2.63	11.54	11.12	6.28				
	PLACE1010624	292.79	228.16	204.46	212.8	278.86	210.51	166.11	171.99	155.01	*		-	-
	PLACE1010628	48.66	44.83	51.05	27.25	26.82	37.41	15.79	19.04	21	*	**	-	-
30	PLACE1010629	19.03	17.43	20.71	12.12	16.38	15.36	14.93	10.66	11.67	*	*	-	-
	PLACE1010630	15.62	16.61	29.46	8.82	19.7	10.88	11.96	14.1	11.42				
	PLACE1010631	22.99	34.46	25.54	20.07	21.82	17.69	19.23	10.89	16.98	*		-	-
	PLACE1010651	8	10.03	11.34	5.4	9.54	5.75	4.35	5.43	5.06	**	*	-	-
	PLACE1010661	13.76	16.34	32.78	7.67	13.64	27.31	6.28	10.55	7.26				
	PLACE1010662	28.68	22.75	30.53	8.67	12.27	9.85	25.45	28.04	24.56	**	*	-	-
35	PLACE1010668	37.33	37.7	40.55	22.62	25.66	20.72	36.78	44.21	37.46	**	*	-	-
	PLACE1010702	12.3	6.26	13.51	6.5	7.73	5.88	8.16	17.08	6.46				
	PLACE1010709	70.65	75.49	68.95	55.09	47.9	57.04	50.07	52.69	55.39	**	**	-	-
	PLACE1010713	80.41	68.48	77.71	36.46	48.82	48.26	28.63	32.14	23.64	**	**	-	-
	PLACE1010714	14.85	12.78	15.24	4.19	7.98	4.28	3.33	8.97	5.69	**	*	-	-
	PLACE1010716	10.07	15.73	17.8	4.88	22.94	9.1	5.99	13.73	3.7				
40	PLACE1010717	16.27	25.02	15.64	13.63	18.64	10.49	10.76	15.16	3.55				
	PLACE1010720	27.49	32.65	34.2	14.65	18.34	8.25	20.45	22.85	22.74	**	*	-	-
	PLACE1010739	32.05	33.62	24.71	15.33	21.42	15.16	17.5	28.68	18.95	*		-	-
	PLACE1010743	44.76	28.81	34.77	15.52	24.11	19.41	14.02	9.67	5.8	*	**	-	-
	PLACE1010752	26.13	19.45	22.02	11.29	9.49	11.37	16.27	24.35	13.97	**	*	-	-
	PLACE1010761	15.13	12.05	32.55	15.98	9.45	8.78	11.14	13.99	12.89				
45	PLACE1010771	28.91	75.96	37.99	25.83	90.51	39.2	31.02	18.86	25.47				
	PLACE1010784	98.45	116.91	93.11	48.56	63.14	52.26	30.32	31.37	33.38	**	**	-	-
	PLACE1010786	16.19	15.53	16.87	8.23	15.52	9.85	7	6.49	9.87	**	*	-	-
	PLACE1010789	46.25	33.26	40.6	14.8	19.17	10.09	44.64	47.22	35.88	**	*	-	-
	PLACE1010800	13.97	9.78	14.67	8.91	6.8	5.82	4.05	5.43	4.76	*	**	-	-
	PLACE1010802	13.35	16.74	27.18	16.21	23.96	4.91	12.32	5.78	37.85				
50	PLACE1010811	26.9	21.58	31.28	13.09	18.01	12.69	11.33	11.86	10.25	*	**	-	-
	PLACE1010813	10.21	6.96	9.67	4.84	4.38	4.87	3.92	2.83	2.64	*	**	-	-
	PLACE1010827	10.79	10.81	12.05	5.43	7.46	5.87	6.65	5.02	3.76	**	**	-	-
	PLACE1010833	9.18	9.98	8.82	3.14	6.94	3.42	1.97	2.07	1.97	*	**	-	-
	PLACE1010839	24.14	30.08	20.18	11.7	20	12.82	16.21	11.13	15.42	*	*	-	-
	PLACE1010856	15.47	10.08	17.88	6.16	6.58	4.26	11.23	8.32	12.26	*		-	-
55	PLACE1010857	32.67	31.48	18.71	21.99	18.34	15.41	16.23	15.82	15.61				
	PLACE1010870	5.83	4.92	4.28	3.87	3.55	2.92	2.91	2.04	1.94	*	**	-	-
	PLACE1010877	26.07	20.6	20.89	14.13	8.42	9.71	13.42	12.71	7.25	**	*	-	-

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	PLACE1009931	29.02	24.07	26.54	12.24	15.5	14.51	9.57	9.79	10.08	**	**	-	-
	PLACE1009935	8.86	7.44	9.7	2.78	11.02	5.42	14.95	3.43	6.35			-	-
5	PLACE1009947	6.29	6.77	9.33	4.25	4.43	3.14	2.04	2.93	4.01	*	*	-	-
	PLACE1009961	7.86	10.21	16.93	5.24	6.64	4.49	4.58	6.45	5			-	-
	PLACE1009971	7.79	7.18	10.26	3.38	7.18	3.34	3.67	4.69	3.08	*	*	-	-
	PLACE1009982	20.35	18.31	15.25	6.6	11.54	11.1	7.87	5.59	13.47	*	*	-	-
	PLACE1009992	12.32	8.01	11.62	6.39	8	6.72	5.24	5.76	6.38	*	*	-	-
	PLACE1009995	21.27	22.11	27.33	13.99	19.4	11.61	8.12	16.7	12.66	*	*	-	-
10	PLACE1009997	23.76	8.03	15.66	6.79	12.9	5.63	10.63	9.44	8.19			-	-
	PLACE1010002	14.56	6.35	12.19	4.66	8.08	5.35	5.32	14.28	7.75			-	-
	PLACE1010011	15.51	14.11	17.86	6.19	7.34	6.29	9.9	6.46	10.43	**	*	-	-
	PLACE1010013	6.16	5.13	12.68	2.37	2.99	1.56	3.53	4.24	2.98			-	-
	PLACE1010021	15.95	20.29	19.12	7.11	14.51	9.31	5.22	8.04	6.15	*	**	-	-
	PLACE1010023	27.49	31.11	21.32	17.41	29.7	20.44	16.49	7.28	6.33	*	*	-	-
15	PLACE1010031	20.87	15.58	32.54	10.16	16.34	7.15	9.85	8.36	10.28			-	-
	PLACE1010039	20.5	6.62	10.22	3.99	6.51	3.25	2.91	6.23	3.6			-	-
	PLACE1010045	11.17	9.8	13.16	5.75	8.43	6.1	5.11	5.8	5.51	*	**	-	-
	PLACE1010053	7.04	5.22	9.04	6.47	4.19	2.39	2.32	2.12	1.16	*	*	-	-
	PLACE1010060	18.9	18.89	15.21	7.4	8.59	8.37	12.03	13.21	10.77	**	*	-	-
	PLACE1010069	10.06	6.9	13.66	5.33	6.93	4.79	2.86	4.42	3.79	*	*	-	-
20	PLACE1010070	8.15	7.09	9.72	2.34	9.93	4.09	3.65	3.87	2.87	**	*	-	-
	PLACE1010074	63.56	56.74	39.11	24.21	42.72	26.72	25.32	15.29	17.22	*	*	-	-
	PLACE1010076	60.58	44.35	55.45	21.22	23.2	19.4	13.41	12.75	17.59	**	**	-	-
	PLACE1010078	13.76	11.23	14.35	6.69	10.2	8.63	4.38	4.87	4.45	*	**	-	-
	PLACE1010081	27.34	19.16	21.57	12.55	15.05	11.39	11.73	7.47	11.77	*	*	-	-
	PLACE1010083	7.1	7.48	5.64	1.97	2.64	2.62	1.69	2.61	1.96	**	**	-	-
25	PLACE1010089	11.95	9.65	9.57	5.5	5.81	4.38	4.84	2.48	4.43	**	**	-	-
	PLACE1010096	14.85	29.75	16.35	5.81	14.35	9.63	7.28	8.67	5.49			-	-
	PLACE1010102	22.29	34.08	23.15	9.26	16.03	13.51	7.63	9.06	7.61	*	**	-	-
	PLACE1010105	18.16	20.35	15.99	6.52	12.73	7.6	4.02	4.82	4.1	*	**	-	-
	PLACE1010106	19.44	22.57	15.73	8.29	8.94	12.84	10.96	10.43	10.44	*	*	-	-
	PLACE1010130	6.82	6.56	4.7	3.25	3.39	2.18	3.26	3.47	3.87	*	*	-	-
30	PLACE1010132	20.18	25.39	20.56	20.18	19.1	12.29	12.71	11.47	12.5	**	**	-	-
	PLACE1010134	10.35	11.42	9.66	6.05	6.05	5.55	6.72	4.12	2.88	**	**	-	-
	PLACE1010139	74.68	88.11	79.85	37.26	51.16	43.84	15.68	26.17	18.6	**	**	-	-
	PLACE1010148	10.48	7.43	9.71	13.14	7.54	4.86	3.27	4.13	7.63			-	-
	PLACE1010156	9.79	7.58	7.63	3.3	5.89	2.96	2.11	2.26	2.53	*	**	-	-
	PLACE1010156	32.59	25.06	24.61	14.7	18.09	11.09	7.29	10.58	14.48	*	**	-	-
35	PLACE1010161	8.48	13.16	10.69	5.15	8.21	6.03	4.47	3.31	5.46	*	*	-	-
	PLACE1010181	8.66	8.08	6.51	2.62	6.54	5.02	3.74	4.12	2.93	**	*	-	-
	PLACE1010194	8.57	7.46	8.67	3.93	4.98	3.84	2.86	2.08	2.37	**	**	-	-
	PLACE1010202	8.39	6.76	12.45	8.31	10.9	5.79	3.56	3.42	2.04	*	*	-	-
	PLACE1010231	12.97	10.31	14.49	7.96	15.61	7.14	8.93	9.42	8.37	*	*	-	-
40	PLACE1010235	12.62	15	11.24	4.59	11.54	4.53	6.11	5.14	4.76	**	*	-	-
	PLACE1010237	5.04	3.77	4.4	2.18	2.37	2.25	2.72	0.64	1.32	**	*	-	-
	PLACE1010251	9.46	4.47	6.29	5.09	4.79	10.14	3.78	5.92	3.88			-	-
	PLACE1010261	8.26	6.41	4.7	3.42	2.86	2.61	4.52	5.73	4.64	*	*	-	-
	PLACE1010270	7.63	8.07	6.36	3.29	6.44	3.19	3.88	4.95	3.8	**	*	-	-
	PLACE1010273	8.97	10.55	5.7	2.72	5.68	3.04	2.78	2.04	3.83	*	*	-	-
45	PLACE1010274	20.97	18.72	29.56	10.88	16.29	14.38	6.73	5.73	6.16	**	*	-	-
	PLACE1010277	15.29	14.99	14.3	8.87	13.93	8.53	11.24	4.25	6.68	*	*	-	-
	PLACE1010293	16.94	20.61	17.07	9.41	11.63	8.24	17.5	8.1	13.23	**	*	-	-
	PLACE1010297	38.08	33.94	34.95	17.91	31.69	15.63	31.8	20.1	25.32			-	-
	PLACE1010300	21.55	15.58	11.87	9.15	8.78	7.88	3.82	5.36	5.08	*	*	-	-
	PLACE1010310	323.02	293.14	231.29	170.36	221.86	136.69	214.56	235.35	207.22	*	*	-	-
50	PLACE1010321	10.7	10.58	7.81	4.96	7.93	6.91	5.6	6.6	6.56	*	*	-	-
	PLACE1010324	6.25	5.69	6.54	2.88	4.51	2.6	4.43	3.34	2.72	*	**	-	-
	PLACE1010329	14.25	11.37	11.93	5.93	10.64	4.89	9.64	9.84	9.32	*	*	-	-
	PLACE1010330	12.28	14.21	10.49	11.84	10.18	7.63	12	7.94	17.86			-	-
	PLACE1010335	27.7	52.66	38.23	18.39	9.7	13.5	16.61	15.38	28.61	*	*	-	-
	PLACE1010341	6.44	4.91	4.72	2	3.7	1.35	4.14	2.93	4.21	*	*	-	-
55	PLACE1010342	2.35	2.93	2.85	2.16	1.61	1.84	1.14	1.19	0.83	*	**	-	-
	PLACE1010346	10.42	6.88	7.23	4.08	5.74	2.77	5.84	6.92	5.88	*	*	-	-
	PLACE1010362	13.25	8.57	9.24	6.56	7.06	3.39	11.5	13.76	14.05			-	-

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	PLACE1009477	13.59	11.11	12.8	10.3	9.83	6.89	8.75	11.15	10.89				
	PLACE1009493	5.63	4.68	4.85	3.04	3.33	2.25	3.06	2.29	1.49	**	**	-	-
5	PLACE1009502	11.43	9.62	8.39	5.92	5.81	5.8	4.63	7.61	3.19	*	*	-	-
	PLACE1009524	7.56	7.26	7.63	1.87	3.86	2.97	2.66	3.37	2.46	**	**	-	-
	PLACE1009527	18.34	12.36	9.71	6.88	11.49	9.95	6.88	5.14	3.52	*	*	-	-
	PLACE1009531	25.38	45.76	31.03	26.02	25.63	22.82	19.07	17.64	15.03				
	PLACE1009535	12.3	11.08	8.3	3.83	6.2	5.84	3.36	3.71	4.06	*	**	-	-
	PLACE1009539	8.83	10.23	7.02	4.34	7.56	5.81	5.39	6.61	6.43				
10	PLACE1009540	37.43	26.63	27.99	14.5	25.74	28.63	21	11.9	8.26	*	*	-	-
	PLACE1009542	12	5.45	5.7	3.62	4.51	6	3.18	3.69	2.67				
	PLACE1009546	12.67	10.42	6.26	3.72	5.36	5.98	4.35	10.8	3.66				
	PLACE1009556	7.91	6.37	6.72	5.1	4.62	3.16	3.16	2.46	2.95	*	**	-	-
	PLACE1009569	11.99	15.44	10.67	4.76	13.77	8.59	5.94	5.33	6.24	**	**	-	-
	PLACE1009571	7.82	9.37	5.59	3.8	7.13	3.69	5.84	2.83	4.48				
15	PLACE1009573	22.09	15.96	12.31	9.76	17.27	9.7	8.26	8.91	7.87	*	*	-	-
	PLACE1009576	13.53	9.49	9.65	4.21	6.31	4.13	5.49	5.78	4.15	*	*	-	-
	PLACE1009580	9.86	9.33	7.56	5.13	7.69	4.5	8.75	3.59	4.85				
	PLACE1009581	12.95	12.1	8.15	7.19	5.84	4.99	3.45	2.48	3.53	*	**	-	-
	PLACE1009587	13.3	7.18	8.26	7.06	6.26	5.41	3.85	4.9	4.28				
	PLACE1009593	18.08	14.97	12.54	14.73	10.76	9.75	5.8	5.66	6.75	**	**	-	-
20	PLACE1009595	24.28	22.3	15.19	12.22	14.76	12.77	19.19	14	13.81				
	PLACE1009596	8.31	5.39	3.71	3.49	5.52	2.24	4.95	2.48	3.3				
	PLACE1009600	19.52	17.07	12.01	6.59	8.79	3.99	8.49	7.24	8.2	*	*	-	-
	PLACE1009604	19.98	10.38	8.96	4.18	6.3	6.58	3.83	5.54	4.55				
	PLACE1009607	17.2	18	14.19	8.35	9.63	7.56	19.54	17.62	14.33	**	**	-	-
	PLACE1009613	8.31	8.44	6.06	5.1	6.18	4.22	4.39	2.55	3.96	*	*	-	-
25	PLACE1009621	18.02	17.88	15.95	7.12	14.76	8.85	8.46	6.45	12.83	*	*	-	-
	PLACE1009622	16.93	9.51	8.44	5	5.84	4.46	4.96	3.18	4.98				
	PLACE1009624	23.04	19.02	15.23	8.31	9.71	5.05	7.94	5.67	7.85	*	**	-	-
	PLACE1009637	8.56	7.93	7.76	3.81	7.55	4.55	3.93	3.27	4.29	**	**	-	-
	PLACE1009639	14.9	6.92	3.57	7.62	5.31	8.34	6.06	5.69	7.84				
	PLACE1009654	23.03	16.88	13.87	7.6	11.88	8.66	7.4	8.09	10.22	*	*	-	-
30	PLACE1009659	17.62	16.71	15.38	8.59	15.54	12.83	6.78	8.01	7.95	**	**	-	-
	PLACE1009665	13.55	12.11	8.08	5.71	10.66	10.87	8.53	10.59	6.01				
	PLACE1009689	10.91	11.58	11.44	5.71	8.38	5.23	8.32	8.02	8.98	**	**	-	-
	PLACE1009670	10	4.03	5.32	3.93	4.76	8.45	5.57	4.18	5.29				
	PLACE1009708	13.41	10.39	7.63	11.97	12.63	3.57	6.84	3.75	9.58				
35	PLACE1009721	15.79	10.82	9.31	5.16	6.52	4.42	9.11	3.24	6.91	*	*	-	-
	PLACE1009731	5.6	6.57	11.06	5.5	8.58	5.78	5.45	4.8	9.13				
	PLACE1009735	9.43	10.36	12.52	8.67	8.77	6.91	8.15	10.58	12.7				
	PLACE1009737	8.36	8.02	10.98	5.74	17.02	9.89	4.98	11.47	4.66				
	PLACE1009741	8.67	7.59	11.34	6.59	7.8	4.16	2.63	4.89	4.5	*	*	-	-
	PLACE1009752	9.51	12.78	18.39	7.46	12.57	11.15	4.36	9.17	6.85				
40	PLACE1009763	16.81	19.39	15.73	12.5	15.24	13.69	4.1	8.02	6.41	**	**	-	-
	PLACE1009766	7.54	8.76	7.16	6.98	11.81	6.16	9.17	7.21	6.61				
	PLACE1009772	12.62	18.28	13.46	16.58	26.84	15.97	7.94	11.57	13.51				
	PLACE1009782	7.96	6.95	7.99	4.24	6.2	9.33	6.26	4.41	4.35	*	*	-	-
	PLACE1009794	8.71	9.98	15.31	6.91	7.94	5.64	5.81	8.68	9.35				
	PLACE1009798	15.7	11.58	19.23	8.55	9.28	7.81	5.76	14.83	6.68	*	*	-	-
45	PLACE1009845	5.69	8.07	10.29	5.79	6.81	9.4	3.27	3.21	4.35	*	*	-	-
	PLACE1009849	4.29	6.11	7.54	5.25	6.21	5.82	3.81	4.56	5.57				
	PLACE1009857	5.97	7.92	9.09	4.47	4.42	3.88	4.36	2.37	3.86	*	*	-	-
	PLACE1009861	21.52	19.65	20.31	10.08	12.87	8.67	13.97	16.84	16.62	**	**	-	-
	PLACE1009872	225.97	496.72	352.07	405.22	548.08	335.22	583.26	839.73	801.36	*	*	-	-
	PLACE1009877	93.23	95.23	121.84	37.51	29.99	31.46	33.83	34.28	50.76	**	**	-	-
50	PLACE1009879	8.31	6.4	8.23	4.6	3.94	4.45	5.4	6.04	5.14	**	**	-	-
	PLACE1009886	6.21	5.71	6.04	3.1	4.43	3.64	3.76	3.74	3.91	**	**	-	-
	PLACE1009888	16.61	15.12	18.96	7.12	11.3	10.4	7.67	15.82	6.68	*	*	-	-
	PLACE1009908	10.69	13.22	16.94	7.9	15.1	13.2	5.89	21.59	11.16				
	PLACE1009919	18.85	20.8	15.18	25.14	44.25	16.53	6.38	10.08	17.42				
	PLACE1009921	12.5	11.43	13.89	6.9	11.45	10.84	6.74	7.53	6.61	**	**	-	-
55	PLACE1009923	37.38	53.14	35.03	28.71	69.72	45.4	23.4	36.13	28.69				
	PLACE1009924	26.03	16.67	21.48	6.94	14.99	12.69	28.73	40.32	10.81				
	PLACE1009926	10.23	25.64	14.3	6.85	14.35	14.92	6.32	35.83	7.94				

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	PLACE1008930	19.58	15.04	12.13	4.18	13.75	5.17	6.22	5.55	4.1	*	-
	PLACE1008934	16.24	5.52	7.38	4.49	6.52	4.37	6.49	7.37	5.19		
5	PLACE1008941	7.73	6.12	5.84	2.96	5.21	1.59	3.04	3.68	3.45	**	-
	PLACE1008947	14.65	13.48	10.43	5.14	6.75	4.89	11.51	10.32	9.84	**	-
	PLACE1008984	13.25	12.98	9.34	5.45	4.62	4.47	6.01	1.98	3.15	**	-
	PLACE1008985	7.07	4.44	3.91	3.08	5.06	1.59	3.53	4.2	7.06	*	-
	PLACE1008994	3.13	3.32	2.46	1.43	1.88	1.44	1.4	0.58	1.95	**	-
	PLACE1009020	10.8	10.64	11.79	9.97	10.93	10.07	6.16	7.97	6.48	**	-
10	PLACE1009027	3.76	1.41	3.27	1.39	2.64	1.03	1.54	3.8	0.71	*	-
	PLACE1009039	4.77	3.08	3.94	1.85	3.67	1.57	1.09	2.8	1.86	*	-
	PLACE1009045	8.82	6.04	9.52	3.29	4.82	2.76	2.8	7.24	2.61	*	-
	PLACE1009048	2.06	2.51	4.63	1.04	3.38	1.38	0.62	1.42	2.18		-
	PLACE1009050	3.79	3.3	4.62	1.78	8.12	1.71	2.85	3.16	1.84		-
15	PLACE1009060	19.41	22.53	16.93	15.31	14.46	12.87	9.32	9.93	14.15	*	-
	PLACE1009067	9.68	7.46	10.13	4.56	10.58	6.42	9.09	6.46	12.36		-
	PLACE1009071	20.58	20.38	20.82	12.02	12.73	10.86	11.69	13.76	13.78	**	-
	PLACE1009090	9.2	7.98	10.33	4.26	8.7	11	5.63	8.05	9.63	*	-
	PLACE1009091	83.22	75.58	81.93	64.69	72.29	69.99	50.11	57.74	52.35	*	-
	PLACE1009094	10.97	9.99	9.32	7.26	14.47	8.52	5.42	12.88	9.03		-
20	PLACE1009099	8.3	10.72	9.62	5.36	7.81	7.04	3.55	4.21	4.97	*	-
	PLACE1009110	7.1	4.33	5.05	4.77	6.82	3.04	3.19	5.04	4.48		-
	PLACE1009111	4.96	7.16	3.67	5.5	7.31	3.03	4.35	3.44	4.4		-
	PLACE1009113	16.47	13.6	10.85	8.16	11.25	17.03	7.41	8	12.76		-
	PLACE1009130	13.26	8.46	18.05	9.61	8.47	7.77	7.71	8.47	5.61		-
	PLACE1009150	9.24	9.08	10.56	5.69	8.31	5.53	5.87	7.76	4.55	*	-
25	PLACE1009155	15.03	18.94	19.94	9.19	13.17	10.27	14.29	28.01	20.9	*	-
	PLACE1009158	11.31	7.28	10.08	9.16	8.65	7.93	4.47	5.84	5.53	*	-
	PLACE1009166	6.72	5.63	9.02	5.97	6.75	3.86	4.53	5.03	5.29		-
	PLACE1009172	12.12	10.58	13.44	7.25	11.43	5.16	4.32	7.13	6.22	**	-
	PLACE1009174	15.11	18.12	12.65	13.74	20.86	11.26	7.87	11.41	11.21		-
	PLACE1009183	26.21	21.05	18.45	14.71	13.12	9.63	13.01	10.13	17.14	*	-
30	PLACE1009186	16.33	12.69	14.24	9.34	9.66	8.31	3.27	14.09	4.74	**	-
	PLACE1009190	10.56	9.79	11.19	7.65	14.06	8.19	4	6.56	7.14	*	-
	PLACE1009196	13.22	10.75	11.8	7.38	9.71	7.53	10.55	12.57	8.91	*	-
	PLACE1009200	19.33	23.87	22.95	10.45	11.73	10.18	10.96	16.99	17.03	**	-
	PLACE1009217	14.13	24.09	16.64	11.71	10.57	10.58	3.82	4.64	6.12	*	-
	PLACE1009230	27.31	26.63	32.64	10.81	26.12	18.34	16.83	20.7	24.31	*	-
35	PLACE1009236	9.03	10.62	12.67	7.18	9.87	8.71	6.77	9.24	7.63		-
	PLACE1009246	21.48	34.44	20.99	9.29	20.22	19.2	9.84	35.09	13.46		-
	PLACE1009265	18.86	26.11	25.51	7.08	18.04	9.4	9.34	41.89	9.73	*	-
	PLACE1009279	13.38	8.53	11.2	5.1	7.71	6.09	11.6	8.18	8.91	*	-
	PLACE1009298	49.41	26.44	34.52	26.24	34.87	27.55	30.84	25.04	29.21		-
	PLACE1009308	9.89	13.34	12.22	3.57	6.53	4.13	6.2	4.88	5.8	**	-
40	PLACE1009319	11.98	14.82	14.36	7.73	14.05	5.85	5.29	9.69	4.23	*	-
	PLACE1009328	12.86	13.58	15.59	6.43	10.45	5.86	7.1	8.23	11.41	*	-
	PLACE1009335	8.34	6.5	7.11	3.99	9.88	4.55	4.35	7.62	2.32		-
	PLACE1009338	14.56	12.99	17.79	6.9	13.74	10.92	6.69	5.83	8.27	**	-
	PLACE1009344	6.92	5.13	5.98	3.02	11.98	4.73	3.18	3.89	2.88	*	-
	PLACE1009355	23.17	22.99	26.45	20.27	18.4	17.42	10.13	10.79	9.83	*	-
45	PLACE1009368	8.63	5.09	6.32	3.35	3.66	2.95	3.35	6.92	2.66	*	-
	PLACE1009375	8.35	5.94	11.71	3.65	4.61	3.52	3.31	4.45	3.97	*	-
	PLACE1009388	18.28	22.47	42.99	8.1	17.38	17.02	11.33	10.99	17.5		-
	PLACE1009398	16.02	15.66	15.45	5.65	12.56	7.13	6.81	5.4	4.18	*	-
	PLACE1009404	17.98	10.01	28.67	4.73	28.94	16.4	9.28	9.88	5.15		-
	PLACE1009410	13.78	15.39	9.6	5.62	8.24	5.55	5.28	4.64	5.63	*	-
50	PLACE1009417	16.73	9.41	10.65	12.79	9.15	6.2	6.71	6.82	7.05	*	-
	PLACE1009424	9.71	6.5	9.08	4.75	4.94	4.28	3.44	4.18	4.16	*	-
	PLACE1009434	21.73	12.22	10.01	6.12	7.87	5.95	3.69	5.83	2.87	*	-
	PLACE1009443	7.33	14.79	8.94	2.98	3.24	2.54	4.85	2.06	4.82	*	-
	PLACE1009444	13.13	14.77	17.33	7.6	10.32	8.01	6.52	5.24	6.76	*	-
	PLACE1009459	8.33	7.91	9.2	3.33	5.72	4.79	3.49	3.56	3.33	**	-
55	PLACE1009460	9.94	14.6	11.32	7.7	8.3	7.62	5.02	3.61	2.95	*	-
	PLACE1009468	12.43	12.85	9.11	5.4	5.98	4.76	4.66	5.65	8.17	**	-
	PLACE1009476	7.02	4.9	5.38	3	3.07	3.32	3.74	3.76	3.61	*	-

Table 486

	PLACE1008426	9.02	4.49	7.91	3	6.94	4.35	2.2	4.69	5.02			
	PLACE1008429	6.72	4.48	8.18	2.61	5.16	2.94	4.14	3.81	3.51			
5	PLACE1008430	14.21	10.67	13.56	6.18	13.27	13.78	7.02	13.13	6.7			
	PLACE1008437	10.05	12.15	13.23	7.3	7.15	7.37	5.37	12.87	6.94	**	-	-
	PLACE1008453	9.93	7.46	12.22	2.24	6.5	5.48	4.46	4.26	3.11	*	*	-
	PLACE1008454	21.6	22.78	23.98	10.1	17.4	10.34	15.68	15.08	11.82	*	**	-
	PLACE1008455	34.62	30.32	25.55	12.02	19.33	14.29	16.61	22	23.57	*	*	-
	PLACE1008457	11.67	8.98	10.52	6.51	8.83	5.84	9.41	9.29	9.88	*		-
10	PLACE1008465	7.78	1.83	9.65	2.4	6.54	3.49	3.66	7.05	5.33			-
	PLACE1008469	9.74	7.14	10.33	5.36	5.23	4.43	4.05	9.23	7.36	*		-
	PLACE1008488	7.69	9.06	16.27	2.87	27.47	11.79	3.3	10.09	3.95			-
	PLACE1008519	13.67	13.58	16.65	12.54	13.83	10.17	7.48	12.69	9.26			-
	PLACE1008524	16.01	17.28	16.34	7.24	7.92	7.96	6.91	6.21	5.65	**	**	-
	PLACE1008531	10.83	9.9	10.84	3.6	7.89	4.88	5.5	5.84	4.3	*	**	-
15	PLACE1008532	31.22	27.65	29.16	13.43	17.17	10.64	20	23.65	23.08	**	**	-
	PLACE1008533	16.83	14.02	15.02	5.15	9.56	6.45	8.04	10.1	5.87	**	**	-
	PLACE1008542	11.05	6.89	12.38	7.62	8.01	5.5	3.52	4.43	4.89	*		-
	PLACE1008549	8.08	8.32	6.56	2.76	4.31	2.64	3.19	4.98	1.09	**	*	-
	PLACE1008560	9.22	6.82	8.11	4.41	4.7	3.86	2.93	3	1.98	**	**	-
	PLACE1008567	12.46	12.8	11.19	3	4.87	3.75	6.88	4.77	3.6	**	**	-
20	PLACE1008568	11.23	14.36	12.98	7.02	10.07	9.2	4.7	4.88	4.71	*	**	-
	PLACE1008569	26.76	24.83	18.12	14.19	21.2	18.33	7.27	6.81	8.32	**		-
	PLACE1008584	12.08	11.9	8.94	4.22	7.86	8.69	3.89	8.17	12.55			-
	PLACE1008585	31.41	26.84	21.37	12.16	16.5	14.29	13.28	12.03	8.95	*	**	-
	PLACE1008603	88.44	58.86	85.85	77.22	107.17	81.02	46.69	50.43	38.76	*		-
	PLACE1008621	8.51	5.24	9.31	3.02	5.85	4.1	5.45	4.49	4.31			-
25	PLACE1008625	5.93	4.92	6	3.54	3.65	3.16	1.59	3.39	1.97	**	**	-
	PLACE1008626	7.17	7.14	8.41	4.47	6	4.22	1.34	2.51	2.04	*	**	-
	PLACE1008627	16.63	17.04	15.47	5.85	10.41	6.49	6.26	6.1	3.78	**	**	-
	PLACE1008629	17.85	11.92	13.81	7.55	20.73	20.07	19.18	21.96	10.11			-
	PLACE1008630	10.34	9.61	9.25	3.39	5.31	3.93	4.34	4.76	3.98	**	**	-
	PLACE1008643	13.74	8.06	11.06	4.37	9.48	6.43	6.52	10.46	6.53			-
30	PLACE1008650	8.08	8.41	6.76	5.59	6.65	6.55	3.49	3.07	2.46	**		-
	PLACE1008657	20.07	16.69	13.65	10.15	11.39	11.63	13.41	12.65	11.83	*		-
	PLACE1008664	16.27	13.14	9.24	7.33	8.26	11.12	3.83	4.48	3.67	*		-
	PLACE1008693	19.24	12.55	11.48	5.9	7.48	8.22	10.23	10.76	7.26	*		-
	PLACE1008696	15.27	18.92	13.52	8.56	14.35	6	9.09	6.63	6.3	**	**	-
35	PLACE1008715	12.07	13.36	10.97	4.79	8.42	4.36	3.79	3.86	5.13	*	**	-
	PLACE1008716	9.95	8.16	6.63	2.27	4.4	4.51	4.19	6.11	4.54	*	*	-
	PLACE1008722	23.06	17.9	16.82	6.63	6.45	7.4	2.92	4.69	5.05	**	**	-
	PLACE1008738	22.32	18.29	21.17	13.47	19.87	5.51	15.63	19.57	23.28			-
	PLACE1008742	84.96	67.74	75.79	53.3	53.65	55.26	60.09	57.16	55.95	*	*	-
	PLACE1008744	16.51	21.18	12.37	7.32	20.16	7.33	4.32	8.49	4.3	*		-
40	PLACE1008748	11.7	8.68	7.89	3.6	3.19	1.95	5.93	2.92	4.46	**	*	-
	PLACE1008757	11.86	10.89	13.3	7.9	13.9	4.82	5.45	7.52	5.79	**		-
	PLACE1008766	12.58	9.83	17.07	7.19	5.62	5.22	3.78	3.59	4.47	*	*	-
	PLACE1008785	10.19	7.64	9.9	4.61	6.58	4.55	5.8	6.94	7.6	*		-
	PLACE1008790	13.83	10.34	7.69	5.61	9.25	7.49	3.5	9.44	5.2			-
	PLACE1008798	38.55	17.87	28.15	8.39	18.4	13.81	20.54	7.92	18.2			-
45	PLACE1008807	7.39	8.08	6.82	3.5	7.58	4.57	3.41	3.26	4.62	**		-
	PLACE1008808	14.69	14.79	11.52	7.99	7.57	5.08	6.28	4.51	3.87	**	**	-
	PLACE1008813	11.75	10.29	9.69	7.89	7.9	5.45	5.57	3.03	5.81	*	**	-
	PLACE1008836	10.67	11.89	10.55	6.13	6.4	5.79	6.93	5.51	7.49	**	**	-
	PLACE1008851	17.38	11.13	11	56.43	9.1	6.6	16.91	8.29	14.77			-
	PLACE1008854	6.2	6.11	5.28	5.3	4.42	2.5	6.46	5.33	5.1			-
50	PLACE1008864	19.27	13.51	11	10.42	10.96	6.46	14.43	10.21	14.23			-
	PLACE1008867	23.06	15	11.22	6.2	9.84	4.86	5.54	7.61	6.44	*		-
	PLACE1008876	176.15	214.21	204.76	126.27	307.85	120.09	159.59	163.82	275.73			-
	PLACE1008887	10.24	8.9	12.49	4.83	6.78	7.76	5.17	4.7	6.53	*	*	-
	PLACE1008902	13.44	7.31	7.6	4.55	6.21	3.19	5.49	3.64	7.36			-
	PLACE1008911	20.57	23.29	15.12	14.15	10.76	7.02	9.39	4.72	9.35	*	*	-
55	PLACE1008917	7.8	5.68	7.21	2.11	3.42	3.23	3.5	4.03	3.26	**	**	-
	PLACE1008920	3.68	3.98	2.25	1.85	2.38	2.78	2.45	2.52	1.03			-
	PLACE1008925	6.03	2.49	3.19	2.51	4.89	2.3	3.22	3.16	2.76			-

Table 485

	PLACE1007950	123	99.57	90.18	75	73.07	58.93	91.95	114.45	65.58	*	-
	PLACE1007954	8.61	7.27	7.95	5.58	7.49	5.54	3.77	4.03	2.97	**	-
5	PLACE1007955	17.64	15.71	13.66	11.51	9.3	9.77	6.99	7.04	4.12	*	-
	PLACE1007956	11.86	9.09	7.66	6.57	9.25	6.9	4.06	4.11	3.44	*	-
	PLACE1007958	9.03	7.99	9.04	2.65	4.9	3.85	2.65	3.75	2.07	**	-
	PLACE1007965	15.3	16.37	14.75	7.93	11.66	10.38	7.23	6.65	5.87	*	-
	PLACE1007969	24.54	27.08	14.83	16.49	16.9	27.64	8.5	10.51	6.83	*	-
	PLACE1007971	14.86	11.23	10.66	4.6	5.77	4.48	5.47	4.42	6.63	**	-
10	PLACE1007990	13.72	7.91	11.82	6.44	6.58	7.05	6.02	6.86	7.23		-
	PLACE1008000	8.41	4.96	5.71	4.66	6.7	5.87	3.52	3.67	3.05		-
	PLACE1008002	8.38	10.22	5.77	6.22	6.73	3.91	3.28	5.34	3.3	*	-
	PLACE1008037	7.47	6.35	4.25	2.21	4.95	4.35	2.35	1.88	2.58	*	-
	PLACE1008044	23.79	15.95	12.5	7.05	19.86	9.73	6.75	10	6.08		-
	PLACE1008046	19.99	21.06	9.24	5.88	14.67	9.04	6.52	7.55	2.12		-
15	PLACE1008080	15.75	26.6	10.27	8.38	25.41	7.91	16.05	14.95	13.99		-
	PLACE1008092	7.67	5.27	5.41	3.04	5.53	3.46	4.86	3.3	3.17		-
	PLACE1008095	16.14	13.84	10.97	4.61	15.55	7.49	3.99	9.83	5.13	*	-
	PLACE1008105	10.75	13.42	9.63	5.46	13.95	7.7	10.03	7.96	6.35		-
	PLACE1008107	6.1	7.51	5.49	4.1	6.68	4.81	2.94	3.48	3.2	**	-
	PLACE1008111	12.94	8.4	7.21	4.85	8.37	6.82	4.56	1.47	2.41	*	-
20	PLACE1008113	48.12	32.36	36.95	30.18	28.53	18.75	11.39	9.55	15.2	**	-
	PLACE1008122	12.52	11.61	10.25	6.59	8.27	7	4.17	3.46	4.27	**	-
	PLACE1008129	12.19	11.22	9.29	6.85	14.96	11.14	8.9	8.46	6.51		-
	PLACE1008132	21.17	14.41	14.91	9.14	12.76	11.94	8.42	8.33	8.23	*	-
	PLACE1008137	10.44	5.69	4.43	4.58	5.26	3.39	2.11	5.99	2.53		-
	PLACE1008174	12.98	8.11	5.85	5.98	7.01	8.19	6.41	5.88	5.58		-
25	PLACE1008177	5.92	6.91	4.83	3.39	4.55	3.28	3.39	1.53	3.61	*	-
	PLACE1008181	12.18	13.31	12.04	4.81	8.68	5.04	4.65	2.16	4.62	**	-
	PLACE1008195	13.39	9.3	9.62	7.44	7.76	6.35	5.1	3.44	5.91	*	-
	PLACE1008198	12.57	11.23	9.83	6.75	6.36	4.49	7.75	4.45	5.55	**	-
	PLACE1008201	10.88	12.64	8.33	6.13	9.66	7.65	6.3	5.05	9.18		-
	PLACE1008209	10.96	9.72	8.1	4.34	4.16	5.71	10.15	7.4	9.7	**	-
30	PLACE1008226	27.87	17.88	18.73	12.29	14.6	8.3	21.53	17.31	27.74		-
	PLACE1008227	19.93	14.99	13.17	9.05	11.55	7.15	13.25	11.51	13.73	*	-
	PLACE1008231	7.21	4.7	4.12	4.59	4.49	3.26	5.49	4.08	3.39		-
	PLACE1008238	12.79	18.51	15.04	5.47	8.57	11.49	6.6	7.97	4.88	*	-
	PLACE1008244	10.53	9.75	8.28	4.31	8.05	4.21	4.37	5.37	5.6	*	-
	PLACE1008249	15.98	17.02	8.54	6.34	10.36	6.43	17.75	4.65	8.03		-
35	PLACE1008266	102.69	124.96	102.56	88.83	113.4	78.29	73.83	64.62	80.68	*	-
	PLACE1008273	14.29	12.39	14.99	6.95	10.06	11.71	9.42	8.24	5.42	*	-
	PLACE1008275	5.26	4.99	6.79	3.64	10.8	3.9	3.08	3.54	1.54	*	-
	PLACE1008280	8.85	6.62	6.82	4.72	8.46	4.99	3.58	6.79	2.88		-
	PLACE1008282	12.89	9.41	16.7	11.11	18.66	13.55	5.47	11.9	7.88		-
40	PLACE1008297	12.14	13.11	10.26	7.3	9.97	7.71	3.51	6.91	7.44	*	-
	PLACE1008303	11.47	15.79	13.39	11.12	14.28	7.92	5.84	6.87	4.96	**	-
	PLACE1008309	4.41	7.26	5.38	6.88	8.29	6.05	3.56	4.2	4.65		-
	PLACE1008315	19.59	15.1	15.12	9.86	9.1	7.96	7.41	8.94	11.54	**	-
	PLACE1008329	9.22	7.92	9.32	5.09	12.42	9.18	10.32	5.48	5.2		-
	PLACE1008330	11.04	9.94	11.21	6.66	7.9	5.74	7.19	13.56	9.01	**	-
45	PLACE1008331	9	15.63	14.68	4.1	11.97	13.67	4.88	7.64	5.55	*	-
	PLACE1008351	12.84	22	18.28	7.61	13.2	12.68	4.26	8.16	3.26	*	-
	PLACE1008356	13.82	13.87	16.44	11.31	18.42	14.21	4.74	8.97	9.76	*	-
	PLACE1008359	10.61	8.67	5.76	4.5	6.41	4.58	3.38	3.42	3.83	*	-
	PLACE1008363	15.16	18.55	16.33	11.79	13.53	6.82	4.87	5.65	5.64	**	-
	PLACE1008369	3.81	2.08	4.52	1.83	1.71	1.5	2.12	2.77	3.86		-
50	PLACE1008392	11.15	8.41	9.64	4.52	7.73	5.42	7.63	11.47	6.48	*	-
	PLACE1008394	11.83	9.98	11.91	7.27	9.44	5.81	12.55	9.98	4.42	*	-
	PLACE1008398	23.83	19.35	38.34	11.94	10.53	17.6	18.05	34.01	14.62		-
	PLACE1008401	9.05	11.72	16.92	8.36	9.24	6.49	5.59	6.88	7.72		-
	PLACE1008402	8.61	13.91	16.2	7.29	15.86	9.77	4.08	3.77	5.17	*	-
	PLACE1008405	93.35	102.54	109	58.1	65.02	51.5	64.76	63.76	63.34	**	-
55	PLACE1008409	9.15	10.83	10.45	7.65	10.56	6.32	5.13	6.5	6.16	**	-
	PLACE1008420	4.6	3.84	4.48	2.08	3.3	3.48	3.03	3.37	1.89	*	-
	PLACE1008424	8.77	7.89	8.59	4.05	4.03	4.43	4.27	11.48	5.37	**	-

Table 484

	PLACE1007507	16.63	14.22	12.4	6.49	10.52	5.63	19.33	11.77	17.21	*	-
	PLACE1007511	13.25	11.24	10.12	3.93	3.98	2.6	3.01	2.43	4.23	**	**
5	PLACE1007513	4.53	2.47	3.69	3.11	1.64	1.97	3.06	2.87	2.04	*	-
	PLACE1007524	49.4	41.73	37.58	21.38	27.05	16.04	48.44	42.69	43.18	*	-
	PLACE1007525	7.48	5.19	4.73	2.29	4.06	1.12	3.74	3.09	4.25	*	-
	PLACE1007537	8.42	7.3	9.61	4.91	5.72	3.83	6.57	5.7	5.93	*	*
	PLACE1007544	5.67	4.7	5.81	1.28	3.36	2.37	4.39	1.96	2.74	*	*
	PLACE1007547	16.52	13.81	13.32	8.64	13.2	6.38	11.36	9.38	8.52	*	*
10	PLACE1007557	22.7	26.33	16.93	8.14	13.6	7.63	24.98	15.24	20.74	*	-
	PLACE1007560	25.9	29.34	29.57	9.23	13.41	19.28	21.02	29.63	20.17	*	-
	PLACE1007565	3.88	2.26	3.15	1.89	4.77	2.57	1.98	2.22	2.43		
	PLACE1007580	4.59	3.09	5.83	2.2	6.97	1.42	1.87	5.45	1.54		
	PLACE1007583	3.85	2.11	3.18	1.64	4.14	4.12	2.02	2.49	2.5		
	PLACE1007591	5.53	6.61	10.65	2.9	4.26	2.95	4.34	4.96	5.37		
15	PLACE1007598	9.38	9.28	9	2.96	15.1	3.88	2.92	7.29	5.52	*	-
	PLACE1007610	3.93	7.09	11.32	4.32	11.34	3.52	4.35	8.81	6.31		
	PLACE1007618	4.85	4.13	5.18	3.66	4.39	5.01	3.62	3.16	4.04	*	-
	PLACE1007621	20.28	8.69	15.56	6.24	12.9	7.98	10.43	10.06	8.81		
	PLACE1007626	35.43	28.05	45.42	25.74	50.46	26.54	38.37	49.51	35.74		
20	PLACE1007632	8.18	6.61	8.24	5.02	13.18	12.74	5.22	7.37	5.53		
	PLACE1007635	17.84	21.72	22.91	16.26	21.82	12.51	7.83	15.31	12.34	*	-
	PLACE1007645	29.46	44.32	50.79	23.79	28.69	29.97	18.24	19.23	26.32	*	-
	PLACE1007649	5.43	7.86	6.71	4.51	9.04	3.18	4.73	2.18	2.74	*	-
	PLACE1007659	23.48	33.01	29.15	18.02	23.66	14.9	25	27.47	23.31		
	PLACE1007669	17.27	10.25	11.99	6.18	6.01	5.07	11.02	11.92	11.97	*	-
	PLACE1007677	15.06	16.35	15.01	8.53	7.7	12.44	11.15	10.52	8.69	*	**
25	PLACE1007688	30.37	21.3	31.84	9.3	9.6	6.97	5.43	7.59	4.3	**	**
	PLACE1007680	16.29	14.9	13.94	8.07	9.45	10.99	5.55	6.43	7.69	**	**
	PLACE1007697	7.22	8.07	8.65	7.42	10.69	12.34	4.08	7.67	5.36		
	PLACE1007702	9.68	16.27	14.18	8.71	12.06	12.61	6.79	7.15	7.48	*	-
	PLACE1007706	14.91	16.84	14.3	10.72	14.81	6.93	5.13	7.31	8.61	**	-
	PLACE1007725	15.22	10.08	11.32	8.01	8.72	6.5	4.92	7.04	8.09	*	-
30	PLACE1007729	9.62	8.04	8.37	6.72	6.7	9.05	5.49	8.22	6.25		
	PLACE1007730	14.54	9.51	13.82	11.08	11.03	10.92	7.95	12.03	8.17		
	PLACE1007737	26.23	17.29	21.46	12.32	15.98	9.58	13.89	16.95	16.37	*	-
	PLACE1007743	15.82	16.04	17.86	8.71	10.59	7.11	6.26	11.61	4.77	**	*
	PLACE1007746	42.92	48.93	37.39	28.78	37.1	35.26	16.78	10.7	15.4	**	-
35	PLACE1007753	14.39	18.43	15.85	13.55	30.64	24.37	6.15	10.27	7.62	**	-
	PLACE1007769	10.21	11.59	12.77	11.76	15.29	7.33	6.11	7.79	6.57	**	-
	PLACE1007780	5.74	2.24	6.02	2.59	3.98	3.4	2.92	4.75	4.26		
	PLACE1007791	7.33	7.57	11.32	6.64	9.26	5.56	5.87	21.8	5.42		
	PLACE1007807	10.4	6.68	8.59	5.19	10.28	4.84	5.47	15.67	6.17		
	PLACE1007810	21.71	5.64	13.76	5.45	11.67	20.68	5.72	5.81	5.21		
40	PLACE1007814	23.92	18.73	27.74	15.47	17.03	13.06	6.08	14.46	7.51	*	*
	PLACE1007828	9.64	11.73	14.79	8.65	13.95	8.98	3.79	6.99	9.71		
	PLACE1007829	21.85	26.65	22.58	10.61	17.88	12.23	16.05	20.8	14.7	*	-
	PLACE1007841	12.3	14.26	16.38	8.52	14.37	8.97	13.64	10.06	9.74		
	PLACE1007842	7.29	10.66	10.16	4.04	8.15	4.63	12.33	6.69	8.73		
	PLACE1007843	8.16	7.8	8.23	4.71	8.29	4.31	4.49	6.71	2.94	*	-
45	PLACE1007845	14.45	11.07	14.52	8.03	11	7.42	8.69	6.18	6.07	*	*
	PLACE1007846	9.52	9.87	9.08	6.82	5.35	6.86	5.9	8.42	7.28	**	*
	PLACE1007848	10.96	7.82	17.21	7.21	6.58	4.31	3.05	6.37	4.05		
	PLACE1007852	38.45	42.72	20.94	14.02	16.83	14.13	8.86	9.78	9.37	*	*
	PLACE1007858	19.55	23.44	22.4	13.38	19.15	12.06	12.86	10.77	11.27	*	**
50	PLACE1007866	106.06	120.7	80.75	100.84	136.45	101.33	115.22	114.23	135.63		
	PLACE1007871	46.64	38.13	36.79	13.91	18.93	11.35	19.19	30.79	21.75	**	*
	PLACE1007877	11.11	8.45	11.03	7.02	8.84	8.93	6.39	7.32	5.13	*	-
	PLACE1007878	37.18	35.47	41.1	20.44	30.52	24.13	17.29	22.96	22.69	*	**
	PLACE1007881	7.08	8.99	9.43	6.21	6.72	5.42	3.38	3.24	2.37	*	**
	PLACE1007885	13.54	9.97	15.34	8.52	12.53	23.99	7.37	7.86	5.04	*	-
	PLACE1007897	9.52	10.47	10.76	6.15	8.22	8.75	3.33	3.79	2.67	*	**
55	PLACE1007908	27.14	31.91	21.03	11.57	16.17	12.14	8.02	9.92	9.32	*	**
	PLACE1007922	13.27	17.29	11.92	8.56	15.72	12.01	8.76	6.61	6.69	*	-
	PLACE1007946	8.9	7.88	9.69	3.7	5.72	2.73	4.49	7.73	5.93	**	-

Table 483

	PLACE1007045	15.78	21.85	22.19	13.09	30.39	13.83	9.78	12.74	10.75	*	-	-
	PLACE1007048	19.3	14.44	24.23	9.41	10.57	11.83	9.91	12.06	15.93	*	-	-
5	PLACE1007053	9.31	5.29	12.35	5.07	3.06	4.56	5.35	25.19	5.41			
	PLACE1007068	8.67	4.77	10.34	5.02	4.28	3.54	2.63	5.02	2.74			
	PLACE1007070	18.55	13.15	15.29	11.03	10.78	10.43	6.96	7.21	5.68	*	**	-
	PLACE1007076	17.03	17.73	24.8	24.48	28.32	20.68	16.51	16.83	13.48			
	PLACE1007077	7.35	10.92	12.26	4.54	5.11	3.6	6.43	5.8	5.66	*	*	-
	PLACE1007081	41.36	35.79	51.18	37.04	46.87	29.31	28.18	20.46	26.31	*	*	-
10	PLACE1007082	14.74	14.27	12.93	7.51	10.12	5.83	13.51	10.86	10.42	*	-	-
	PLACE1007092	9.52	6.79	8.77	4.99	5.23	4.04	4.42	5.84	7.46	*	-	-
	PLACE1007096	8.7	7.04	11.33	3.9	6.35	2.61	4.31	10.62	7.33	*	-	-
	PLACE1007097	5.03	2.47	6.96	1.99	2.48	1.11	1.28	4.01	1.45			
	PLACE1007099	14.43	9.17	8.68	4.57	6.07	4.14	4.7	5.17	7.04	*	-	-
	PLACE1007105	9.93	9.89	19.72	4.22	5.46	3.96	5.83	6.27	9.12			
15	PLACE1007108	7.87	5.2	13.72	3	3.64	1.32	1.96	5.82	6.69			
	PLACE1007111	24.26	33.98	33.96	8.75	19.56	14.7	10.61	6.46	7.98	*	**	-
	PLACE1007112	13.01	15.69	10.39	7.41	17.53	7.08	9.51	4.37	5.03	*	-	-
	PLACE1007130	7.28	5.04	7.51	3.97	3.76	4.18	4.15	3.12	4.45	*	*	-
	PLACE1007132	11.01	5.03	12.54	6.14	8.23	7.16	4.79	4.84	3.32			
	PLACE1007140	8.1	6.86	9.35	5.74	3.15	5.4	5.1	3.71	4.08	*	*	-
20	PLACE1007143	7.85	4.78	7.3	2.76	4.42	3.52	3.05	2.29	1.7	*	*	-
	PLACE1007189	21.25	20.42	20.37	9.68	16.47	8.01	13.15	12.99	9.63	*	**	-
	PLACE1007178	5.69	6.23	10.18	2.7	4.22	2.51	2.33	2.04	2.65	*	*	-
	PLACE1007190	9.72	11.86	11.22	5.23	6.85	7.11	6.69	5.59	16.1	**	-	-
	PLACE1007201	11.31	9.93	8.97	4.82	6.82	3.46	5.63	4.72	7.11	*	*	-
25	PLACE1007202	14.96	8.93	16.56	5.53	5.66	4.16	5.42	6.71	5.73	*	*	-
	PLACE1007226	110.77	93.69	99.41	39.5	91.72	72.21	82.39	66.08	68.98	*	-	-
	PLACE1007236	53.4	44.43	54.82	32.05	30.06	27.04	24.59	23.68	27.03	**	**	-
	PLACE1007239	22.01	21.57	16.94	9.88	9.77	7.42	7.13	6.76	5.92	**	**	-
	PLACE1007242	8.39	6.41	6.58	2.97	3.53	2.45	2.76	1.3	1.2	**	**	-
	PLACE1007243	51.6	47.43	46.65	38.48	53.48	36.76	30.03	24.3	28.99	**	-	-
30	PLACE1007247	13.13	15.22	14.84	6.24	10.32	8.42	5.62	3.59	3.12	*	**	-
	PLACE1007257	10	9.05	8.75	2.44	7.1	2.96	4.72	2.48	2.64	*	**	-
	PLACE1007274	7.6	6.25	7.77	4.39	3.79	3.63	3.01	2.85	2.82	**	**	-
	PLACE1007276	4.51	6.36	6.4	3.01	2.72	2.17	2.35	3.34	6.03	**	-	-
	PLACE1007282	6.97	4.05	5.61	2.64	2.11	3.56	3.82	3.22	3.39	*	-	-
	PLACE1007286	11.57	9.64	10.32	4.04	6.1	5.17	5.17	3.51	3.72	**	**	-
35	PLACE1007296	16.68	11.68	9.05	4.46	6.87	6.7	3.08	3.72	1.93	*	-	-
	PLACE1007301	14.64	13.62	12.24	3.61	5.07	2.97	5.09	2.86	3.16	**	**	-
	PLACE1007314	9.78	10.14	10.9	5.59	4.99	2.5	3.88	2.16	3.2	**	**	-
	PLACE1007317	8.02	7.14	6.45	3.78	5.35	2.25	2.6	5.13	4.46	*	*	-
	PLACE1007329	6.93	6.45	9.13	3.46	3.47	2.66	3.68	2.3	2.77	**	**	-
	PLACE1007338	57.3	54.81	50.58	25.62	27.62	24.63	49.58	43.5	58.48	**	-	-
40	PLACE1007342	41.9	26.92	32.51	26.5	30.16	25.58	24.13	20.5	25.41			
	PLACE1007345	5.67	4.07	4.44	3.74	2.2	1.81	1.89	2.39	3.76	*	*	-
	PLACE1007346	20.5	15.07	16.53	8.68	6.27	3.82	4.58	3.48	4.29	**	**	-
	PLACE1007359	41.26	41.58	38.73	21.16	26.14	23.96	27.05	27.12	33.74	**	**	-
	PLACE1007367	8.65	8.31	7.96	1.51	3.49	1.31	3.83	1.42	3.11	**	**	-
	PLACE1007375	12.55	10.83	9.7	1.39	5.63	1.68	2.54	2.85	2.14	**	**	-
45	PLACE1007377	6.98	4.2	6.44	4.26	4.62	2.93	3.59	4.11	3.58			
	PLACE1007386	12.07	10.56	14.15	4.68	4.55	5.21	6.12	7.62	8.27	**	*	-
	PLACE1007392	24.21	18.68	19.13	13.92	12.02	7.51	28.83	13.55	12.42	*	-	-
	PLACE1007402	16.76	19.32	15.87	8.17	8.78	7.34	11.49	10.03	10.79	**	**	-
	PLACE1007409	9.51	9.82	11.04	5.11	5.51	7.23	4.08	2.01	3.77	**	**	-
	PLACE1007416	22.54	28.88	22.62	18.84	13.11	6.16	9.22	5.36	7.49	*	**	-
50	PLACE1007420	14.67	16.22	16.74	8.73	9.19	6.79	10.55	5.47	7.91	**	**	-
	PLACE1007431	28.03	25.88	21.18	15.73	16.55	11.3	26.89	15.55	26.27	*	-	-
	PLACE1007450	6.82	6.06	4.68	4.06	4.16	1.71	5.65	4.38	4.35			
	PLACE1007452	14.4	7.71	4.99	5.04	6.12	1.89	4.23	8.8	6.06			
	PLACE1007454	49.36	46.08	36.43	21.1	38.54	20.65	11.86	16.68	14.49	**	-	-
	PLACE1007460	11.52	10.93	12.45	8.16	8.35	4.82	9.8	9.12	12.38	*	-	-
55	PLACE1007470	12.6	13.38	10.86	4.17	8.16	7.05	7.4	4.59	6.97	*	**	-
	PLACE1007484	10.71	6.45	6.68	3.65	4	2.5	4.9	2.97	5.47	*	-	-
	PLACE1007488	14.59	14.99	17.16	7.24	4.48	5.55	6.08	3.67	8.77	**	**	-

Table 482

	PLACE1006610	9.23	9.13	6.93	3.56	3.78	2.24	3.21	1.79	1.8	**	**	-	-
	PLACE1006615	5.13	4.58	7.74	3.64	4.24	3.19	3.42	3.13	4.2			-	-
5	PLACE1006617	10.84	9.51	10.64	5.44	6.44	4.54	8.05	7.55	8.94	**	*	-	-
	PLACE1006618	16.8	17.21	17.66	12.4	9.99	12.65	9.96	9.85	12.85	**	**	-	-
	PLACE1006626	9.9	9.38	11.9	4.89	3.64	3.15	3.8	3.9	3.22	**	**	-	-
	PLACE1006629	97.87	77.5	62.9	75.16	89.13	71.45	50.39	46.85	45	*		-	-
	PLACE1006637	20.62	30.53	27.31	15.03	24.3	12.77	12.77	11.41	9.09	**		-	-
	PLACE1006640	146.09	116.55	121.62	98.73	144.79	100.95	65.85	61.12	81.72	**		-	-
10	PLACE1006644	10.36	13.76	11.01	6.08	7.62	8.85	6.47	3.27	4.59	*	**	-	-
	PLACE1006657	9.47	8.7	8.11	3.37	3.88	4.59	4.46	4.41	6.15	**	**	-	-
	PLACE1006673	128.37	101.12	117.47	60.38	47.28	69.57	78.35	78.65	91.83	**	*	-	-
	PLACE1006678	13.14	7.54	6.57	3.78	6.24	6.14	3.95	3.22	3.7			-	-
	PLACE1006682	5.12	3.88	4.9	2.21	2.14	1.84	2.5	1.43	2.09	**	**	-	-
	PLACE1006684	8.48	6.37	7.5	1.9	2.47	2.03	3.67	2.97	2.03	**	**	-	-
15	PLACE1006688	24.65	21.3	17.88	6.94	8.35	11.27	7.31	3.9	5.72	**	**	-	-
	PLACE1006704	10.75	10.1	9.67	7.04	6.45	4.58	4.88	2.24	3.52	**	**	-	-
	PLACE1006708	6.82	6.62	4.54	1.43	4.58	2.49	4.44	1.95	2.6	*		-	-
	PLACE1006711	7.97	5.26	8.07	5.12	5.44	5.92	5.18	4.48	3.42			-	-
	PLACE1006714	19.78	16.32	14.11	6.39	7.42	10.52	4.46	5.02	4.69	*	**	-	-
	PLACE1006716	21.34	19.47	16.01	5.39	7.72	5.97	12.01	9.74	13.28	**	*	-	-
20	PLACE1006731	8.76	6.59	9.88	4.19	4.28	3.45	3.4	4.54	3.95	*	*	-	-
	PLACE1006754	12.83	10.26	11.39	6.83	5.2	6.24	8.13	4.13	6.6	**	*	-	-
	PLACE1006760	17.66	24.39	15.04	22.5	23.46	11.75	21.46	9.03	14.39			-	-
	PLACE1006779	14.53	14.99	8.23	6.82	4.21	3.38	5.07	3.29	2.36	*	*	-	-
	PLACE1006782	13.34	14.85	8.6	8.19	13.03	7.71	10.87	6.95	6.98			-	-
	PLACE1006783	26.1	20.43	20.94	18.8	23.33	15.04	25.48	18.52	26.77			-	-
25	PLACE1006786	16.5	11.01	11.97	10.81	6.14	7.77	7.76	8.87	8.16	*		-	-
	PLACE1006792	43	28.6	28.89	12.14	32.07	8.88	14.31	11.42	11.6	*	*	-	-
	PLACE1006795	8.46	5.54	6.42	3.27	4.53	2.37	5.98	1.9	3.55	*		-	-
	PLACE1006800	14.69	12.26	14.82	7.42	7.86	5.36	11.02	8.63	12.6	**		-	-
	PLACE1006805	297.4	379.55	330.72	251.58	242.8	138.41	164.1	78.29	121.42	*	**	-	-
30	PLACE1006809	17.57	15.02	14.01	6.89	5.54	5.55	12.59	5.67	9	**	*	-	-
	PLACE1006815	36.78	37.15	24.97	22.04	27.53	19.08	12.45	10.62	14.81	**	**	-	-
	PLACE1006819	8.55	8.96	7.82	3.54	3.27	1.58	3.56	2.07	3.34	**	**	-	-
	PLACE1006820	11.34	11.23	10.69	7.26	7.6	3.88	5.72	6.8	7.48	*	**	-	-
	PLACE1006826	11.68	10.21	8.37	4.26	6.11	3.55	4	5.42	6.14	*	*	-	-
	PLACE1006829	8.66	7.12	6.95	4.95	4.5	3.35	3.73	4.63	3.79	*	**	-	-
	PLACE1006853	7.76	9.04	10.06	4.43	5.8	3.94	5.55	3.81	3.65	**	**	-	-
35	PLACE1006860	13.3	14.13	11.82	7.29	7.57	5.64	7.39	5.43	7.17	**	**	-	-
	PLACE1006867	20.52	19.65	17.03	7.58	12.2	4.52	8.89	4.5	9.57	*	**	-	-
	PLACE1006875	6.86	5.29	3.98	1.69	3.65	1.95	4.64	3.37	3.97	*		-	-
	PLACE1006878	5.96	7.5	9.4	3.45	4.02	4.56	4.29	5.32	7.08	*		-	-
	PLACE1006883	17.58	24.05	25.14	11.43	14.88	9.68	8.41	13.21	13.15	*	*	-	-
40	PLACE1006898	34.11	35.72	38.43	15.89	26.64	17.27	29.81	38.71	32.24	*		-	-
	PLACE1006901	35.59	25.34	29.52	17.57	24.59	23.14	7.06	14.35	10.26	**		-	-
	PLACE1006904	283.91	237.86	265.1	210.53	264.05	203.01	176.62	324.88	215.9			-	-
	PLACE1006917	8.87	8.87	10.42	4.77	9.67	4.27	2.96	3.43	5.44	**		-	-
	PLACE1006932	13.9	19.05	15.16	17.43	14.76	12.47	12.23	11.92	6.34			-	-
	PLACE1006935	5.15	5.68	5.5	4.25	5.6	3.94	4.66	3.59	4.94			-	-
45	PLACE1006956	73	65.43	79.54	37.1	42.04	37.7	45.33	35.06	47.75	**	**	-	-
	PLACE1006958	14.71	17.1	17.16	7.75	10.57	5.67	10.62	14.42	12.92	**		-	-
	PLACE1006959	6.43	6.88	7.06	3.86	5.4	3.09	3.53	7.7	6.81	*		-	-
	PLACE1006961	11.78	13.88	26.99	6.58	7.47	6.43	7.89	10.93	8.94			-	-
	PLACE1006962	8.5	7.71	10.43	6.5	8.54	6.61	4.94	6	6.79	*		-	-
	PLACE1006966	30.69	63.35	44.81	30.7	31	25.61	17.51	20.36	25.7			-	-
50	PLACE1006979	9.85	10.83	9.97	5.88	6.1	5.13	5.32	5.52	6.47	**	**	-	-
	PLACE1006989	23.14	43.35	32.93	27.05	47.4	17.28	23.49	21.59	16.62			-	-
	PLACE1007001	14.41	12.66	17.15	10.97	5.54	6.74	6.74	7.67	8.22	*	**	-	-
	PLACE1007014	9.24	9.35	10.83	6.4	6.04	5.94	4.96	6.16	5.52	**	**	-	-
	PLACE1007021	56.53	38.34	63.46	20.03	40.71	38.97	20.72	29.77	27.88	*		-	-
	PLACE1007026	65.54	56.8	77.56	57.55	60.7	57.93	57.5	58.64	67.29			-	-
55	PLACE1007028	8.5	10.76	17.58	7.89	7.62	6.26	3.31	5.38	3.91	*		-	-
	PLACE1007038	7.96	7.89	13.66	4.19	6.33	3.79	2.05	3.52	4.03	*		-	-
	PLACE1007040	8.13	9.51	15.11	4.74	7.09	4.87	8.84	8.35	6.98			-	-

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	PLACE1006211	18.15	19.46	15.85	7.8	6.56	8.15	10.56	8.16	8.98	**	**	-	-
	PLACE1006219	29.22	24.03	22.82	13.99	7.2	9.8	23.91	20.19	22.03	**	*	-	-
5	PLACE1006223	33.12	26.98	22.85	14.69	11.86	8.81	23.65	22.74	28.77	**	*	-	-
	PLACE1006225	9.38	7.86	6.1	4.18	4.91	3.81	6.34	5.77	7.67	*	*	-	-
	PLACE1006236	23.05	26.18	25.58	13.5	14.02	10.31	12.5	8.37	11.47	**	**	-	-
	PLACE1006238	7.77	5.03	5	2.21	3.14	2.06	2.98	3.13	3.81	*	*	-	-
	PLACE1006245	11.97	10.97	10.73	4.72	5.26	4.25	5.6	4.62	4.52	**	**	-	-
	PLACE1006246	14.07	10.52	9.12	4.17	12.12	4.44	10.44	8.15	11.84		*	-	-
10	PLACE1006248	10.3	9.17	8.56	6.54	7.06	2.85	7.67	4.65	5.48		*	-	-
	PLACE1006262	19.08	26.43	25.66	10.77	26.8	17.9	14.07	10.91	15.08		*	-	-
	PLACE1006269	6.36	4.8	5.62	2.78	2.48	2.42	4.23	3.67	2.13	**	*	-	-
	PLACE1006275	7.84	6.83	8.11	7.79	9.46	8.46	5.82	11.38	4.94		*	-	-
	PLACE1006277	18.16	15.2	14.15	12.95	19.65	15.77	6.69	13.62	7.81		*	-	-
	PLACE1006288	23.16	25	24.73	14.26	21	15.27	15.12	19.66	20.01	*	*	-	-
15	PLACE1006290	3.79	4.35	4.22	1.07	3.07	1.54	0.76	2.83	1.89	*	*	-	-
	PLACE1006298	31.99	33.57	40.35	21.62	22.17	18.83	30.18	26.42	31.31	**	*	-	-
	PLACE1006311	24.06	25.54	26.15	17.19	20.35	18.67	6.49	6.41	12.35	**	**	-	-
	PLACE1006318	23.92	22.31	22.74	9.17	11.04	10.73	16.57	17.39	15.35	**	**	-	-
	PLACE1006325	10.97	6.18	13.93	5.64	6.43	6.48	6.34	4.38	3.68		*	-	-
	PLACE1006331	322.49	249.29	312.14	284.34	344.95	317.78	261.41	376.24	167.86		*	-	-
20	PLACE1006335	6.01	6.97	9.42	6.43	10.77	6.27	4.79	4.63	3.75		*	-	-
	PLACE1006357	23.6	23.65	28.41	15.45	19.92	11.71	10.11	13.97	13.07	*	**	-	-
	PLACE1006360	39.03	55.89	60.83	32.56	29.33	36.42	18.81	22.42	24.3		*	-	-
	PLACE1006384	4.49	3.79	4.52	5.2	5.32	3.04	3.31	4.72	3.85		*	-	-
	PLACE1006385	8.87	13.33	7.94	4.35	7.31	4.11	4.93	6.19	6.08		*	-	-
25	PLACE1006388	7.6	5	7.22	4.35	3.51	2.72	3.12	4.28	3.68	*	*	-	-
	PLACE1006371	12.84	8.89	13.13	6.85	5.51	3.89	5.93	6.19	7.36	*	*	-	-
	PLACE1006373	31.88	31.57	27.89	11.44	15.9	12.05	24.23	17.9	24.85	**	*	-	-
	PLACE1006382	15.81	18.11	17.64	20.83	24.97	21.9	28.68	49.67	48.66	*	*	+	+
	PLACE1006385	23.74	24.67	19.54	18.57	28.47	23.59	6.72	12.92	8.24	**	*	-	-
	PLACE1006391	7.09	8.24	10.32	5.76	7.9	5.35	7.26	7.31	6.16		*	-	-
30	PLACE1006412	14.35	15.98	15.55	14.85	16.87	8.6	8.42	8.73	11.54	**	*	-	-
	PLACE1006414	30.64	63.89	32.69	29.17	49.53	18.72	7.89	13.37	10.44	*	*	-	-
	PLACE1006419	23.86	16.38	19.21	8.34	10.82	5.22	6.84	8.36	7.76	*	**	-	-
	PLACE1006438	16.88	12.75	13.77	5.93	7.83	5.98	6.47	9.15	5.41	**	*	-	-
	PLACE1006443	7.62	8.28	7.5	4.32	3.07	4.42	2.72	3.77	2.68	**	**	-	-
	PLACE1006445	53.01	52.47	65.34	48.09	51.7	50.7	24.1	26.92	30.98	**	*	-	-
35	PLACE1006447	16.45	17.73	17.35	9.64	12.61	9.03	5.67	7.16	5.82	**	**	-	-
	PLACE1006466	5.01	5.76	10.4	3.6	5.26	3.03	3.62	2.47	3.82		*	-	-
	PLACE1006469	9.97	6.93	12.7	4.45	5.94	2.99	6.14	5.96	3.89	*	*	-	-
	PLACE1006470	14.91	13.1	13.03	5.44	14.24	9.05	12.65	15.12	16.52		*	-	-
	PLACE1006472	32.58	24.9	52.53	14.86	22.47	15.58	27.13	60.66	18.13		*	-	-
	PLACE1006476	75.24	88.04	95	49.93	69.66	43.58	73.04	95.42	77.62	*	*	-	-
40	PLACE1006482	19.52	14.87	22.77	7.46	8.75	5.8	9.29	24.88	15.09	**	*	-	-
	PLACE1006488	13.42	9.61	14.11	5.93	5.56	7.22	5.43	7.77	6	*	*	-	-
	PLACE1006492	53.41	60.91	65.09	29.78	38.14	28.4	41.31	52.42	43.57	**	*	-	-
	PLACE1006506	8.93	8.04	10.86	3.97	6.16	3.99	2.77	5.01	4.34	*	**	-	-
	PLACE1006515	6.57	6.17	10.41	2.69	4.8	3.13	3.49	3.57	2.51	*	*	-	-
	PLACE1006516	7.77	7.4	8.18	2.65	6.76	3.8	4.8	4.81	1.69	*	*	-	-
45	PLACE1006520	85.74	60.72	84.5	67.76	73.09	68.07	40.48	30.38	37.31	**	*	-	-
	PLACE1006521	5.4	3.13	7.19	3.67	3.58	2.13	2.56	2.69	1.53		*	-	-
	PLACE1006529	21.37	16.93	24.45	11.55	12.23	11.32	13.1	15.15	16.91	*	*	-	-
	PLACE1006531	8.98	4.63	8.46	3.15	2.87	3.1	2.57	4.69	2.69	*	*	-	-
	PLACE1006534	11.34	12.38	19.26	6.83	8.12	7.2	8.08	5.98	8.82		*	-	-
	PLACE1006540	17.3	13.48	19.89	6	9.01	6.33	5.04	4.77	5.56	**	**	-	-
50	PLACE1006549	57.43	76.29	59.13	33.67	55.25	39.07	49.85	36.47	56.34		*	-	-
	PLACE1006550	27.31	23.34	17.69	10.6	24.86	11.76	7.55	11.98	6.46		*	-	-
	PLACE1006552	41.43	35.42	38.21	18.37	17.9	12.42	28.67	34.48	34.92	**	*	-	-
	PLACE1006557	58.91	37.26	57.05	36.71	38.26	34.22	31.47	33.21	28.65		*	-	-
	PLACE1006563	12.53	9.16	14.73	7.21	10.95	7.06	5.43	4.33	7.28	*	*	-	-
	PLACE1006579	8.16	5.5	6.78	3.56	3.74	3.1	2.74	1.94	3.28	*	**	-	-
55	PLACE1006594	8.47	4.76	7.36	2.29	3.19	1.42	2.51	1.45	1.95	*	*	-	-
	PLACE1006598	14.33	13.22	15.98	9.86	13.18	9.6	7.97	4.5	6.78	**	*	-	-
	PLACE1006607	18.95	13.98	15.61	10.13	11.36	9.36	8.88	8.34	9.06	*	**	-	-

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	PLACE1005828	21.32	17.86	17.66	18.38	28.15	11	8.41	12.77	17.42				
	PLACE1005833	19.04	15.23	17.55	13.68	13.86	12.16	11.62	13.97	12.02	*	*	-	-
5	PLACE1005834	31.23	40.41	42.31	19.53	15.14	12.45	22.97	22.37	27.18	**	*	-	-
	PLACE1005835	8.38	4.15	8.95	3.69	3.05	3.25	2.72	13.83	4.26				
	PLACE1005836	6.34	4.47	-7.11	-2.95	2.55	2.1	1.11	4.41	2.67	*		-	-
	PLACE1005845	7.96	5.73	10.94	3.26	4.7	1.75	3.2	4.47	3.62	*	*	-	-
	PLACE1005850	7.12	8.63	11.96	4.89	4.51	3.39	2.96	4.97	6.83	*		-	-
	PLACE1005851	49.92	45.46	36.8	28.76	36.45	21.25	9.27	9.06	15.36		**	-	-
10	PLACE1005856	9.13	9.45	9.68	4.42	6.52	4.26	5.66	4.96	7.88	**	*	-	-
	PLACE1005875	19.11	12.96	14.66	10.98	11.89	11.34	7.25	9.07	9.2		*	-	-
	PLACE1005876	6.84	5.72	8.26	3.52	3.77	3.26	2.47	4.32	3	*	*	-	-
	PLACE1005878	6.18	5.12	8.15	3.65	4.35	1.9	2.1	7.39	4.67				
	PLACE1005880	10.13	10.1	15.44	15.29	7.51	5.59	11.17	12.56	8.92				
	PLACE1005884	15.13	11.77	15.14	4.99	6.58	3.8	5.03	3.76	7.31	**	**	-	-
15	PLACE1005890	14.12	12.54	15.15	4.55	6.88	4.96	4.84	4.65	10.58	**	*	-	-
	PLACE1005898	22.47	43.11	28.59	9.23	15.21	13.1	10.14	6.45	7.84	*	*	-	-
	PLACE1005913	11.08	12.76	18.39	6.03	9.77	5.43	8.89	7.87	5.55				
	PLACE1005921	38.7	26.83	45.11	14.95	20.53	13.93	14.46	15.3	19.69	*	*	-	-
	PLACE1005923	35.55	28.6	35.14	8.99	9.9	10.33	7.55	9.92	9.16	**	**	-	-
	PLACE1005925	9.28	6.52	9.69	4.22	6.79	3.62	4.36	4.97	3.85	*		-	-
20	PLACE1005927	8.32	7.66	16.91	2.68	3.86	2.87	2.78	4.08	4.5				
	PLACE1005932	17.99	14.04	15.58	5.88	10.67	7.85	11.54	8.65	8.07	*	*	-	-
	PLACE1005934	7.56	7.52	8.85	2.45	3.18	2.78	2.65	3.47	4.68	**	**	-	-
	PLACE1005936	8.91	7.48	10.96	2.6	4	2.78	2.64	2.57	2.62	**	**	-	-
	PLACE1005939	14.59	14.49	13.88	5.4	8.87	6.89	7.24	6.14	6.85	**	**	-	-
	PLACE1005961	184.11	138.22	202.24	63.95	124.2	100.53	163.66	124.76	76.11	*		-	-
25	PLACE1005953	4.12	2.98	5.89	2.81	3.65	1.89	1.88	2.73	1.8				
	PLACE1005955	147.21	90.51	163.74	157.35	162.66	147.78	83.65	73.22	67.87				
	PLACE1005965	21.58	17.57	18.88	12.22	12.03	11.48	18.97	16.95	19.33	**		-	-
	PLACE1005968	7.85	6.96	6.59	3.18	3.12	3.27	2.21	3.25	2.08	**	**	-	-
	PLACE1005975	6.58	6.62	8.7	2.59	2.3	2.44	2.13	2.22	2.24	**	**	-	-
	PLACE1005990	13.31	13.53	10.46	5.9	8.19	4.45	5.05	6.3	5.08	*	**	-	-
30	PLACE1005997	11.27	8.34	8.61	3.34	7.86	4.32	3.83	4.17	2.65		**	-	-
	PLACE1006002	7.2	9.3	8.62	4.13	3.01	2.96	3.03	3.74	3.08	**	**	-	-
	PLACE1006003	19.13	15.27	13.68	6.84	9.41	7.04	14.37	14.53	15.21	*		-	-
	PLACE1006011	14.53	12.13	13.55	5.17	8.37	6.39	12.92	13.32	12.8	**		-	-
	PLACE1006017	17.66	15.33	13.64	7.8	6.85	7.26	3.44	3.4	5.68	**	**	-	-
35	PLACE1006037	19.54	15.64	16.94	5.06	9.69	7.93	15.52	15.45	11.65	**		-	-
	PLACE1006040	21.73	18.19	20.02	8.22	10.07	7.19	9.12	8.1	8.15	**	**	-	-
	PLACE1006063	18.04	22.51	29.29	15.12	25.84	15.72	12.74	9.95	9.64	*		-	-
	PLACE1006071	24.45	22.47	19.59	9.7	14.01	7.6	15.44	10.65	16.68	**	*	-	-
	PLACE1006073	38.01	25.19	30.05	16.58	19.75	14.04	20.88	12.98	14.52	*	*	-	-
	PLACE1006074	21.69	19.68	21.99	13.65	9.89	8.37	10.34	8.49	11.16	**	**	-	-
40	PLACE1006076	6.98	23.32	28.44	3.28	24.98	6.58	8.27	11.5	12.4				
	PLACE1006079	5.36	7.59	6.02	3.43	3.4	2.8	2.82	1.59	5.77	*		-	-
	PLACE1006093	17.29	20.55	15.5	11.28	11.48	8.91	6.51	3.34	7.24	*	**	-	-
	PLACE1006116	6.66	4.88	6.43	2.05	2.48	1.97	3.15	0.18	1.1	**	*	-	-
	PLACE1006119	9.01	6.23	8.79	2.16	3.34	2.75	3.83	1.45	4.03	**	*	-	-
	PLACE1006129	11.8	19.99	13.49	6.74	8.91	6.37	6.93	6.6	9.52	*	*	-	-
45	PLACE1006139	11.03	9.2	11	5.5	4.05	3.54	5.37	5.2	8.18	**	*	-	-
	PLACE1006143	34.11	32.02	23.21	13.95	17.22	16.38	5.72	10.56	8.1	*	**	-	-
	PLACE1006157	20.08	16.11	21.21	6.21	8	10.58	9.26	6.55	9.67	**	**	-	-
	PLACE1006159	22.74	19.17	16.53	5.42	7.39	5.18	11.09	3.24	9.93	**	*	-	-
	PLACE1006164	10.4	12.06	12.25	3.81	6.89	2.98	4.98	2.08	3.84	**	**	-	-
	PLACE1006167	150.43	130.54	128	105.79	82.45	55.54	118.18	93.17	93.5	*	*	-	-
50	PLACE1006170	12.04	9.61	11.88	3.48	3.73	3.68	2.75	2.82	3.04	**	**	-	-
	PLACE1006181	14.32	10.9	10.36	5.74	6.23	4.42	6.29	2.91	4.76	**	*	-	-
	PLACE1006187	23.98	21.9	19.46	7.26	12.29	5.92	18.37	17.04	21.06	**		-	-
	PLACE1006195	27.43	19.87	20.2	13.18	6.92	6.66	10.17	6.16	5.89	*	**	-	-
	PLACE1006196	14.88	10.51	8.01	6.22	7.31	2.89	4.66	4.93	4.37	*		-	-
	PLACE1006197	25.54	27.48	23.66	11.8	19.78	16.24	17.7	18.29	21.21	*	*	-	-
55	PLACE1006198	12.15	16.37	13.46	4.05	5.5	3.12	5.11	3.17	4.8	**	**	-	-
	PLACE1006205	8.36	7.44	9.1	4.62	5.35	3.49	12.53	5	7.53	**		-	-
	PLACE1006208	10.95	10.22	9.53	5.04	9.25	5.28	4.47	4.13	4.33	**		-	-

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	PLACE1005495	11.05	11.48	9.86	5.01	5.38	4.26	3.3	3.97	3.58	**	**	-	-
	PLACE1005497	9.89	9.26	8.02	3.87	6.07	3.47	3.99	3.58	2.73	**	**	-	-
5	PLACE1005499	23.46	21.47	29.66	8.65	19.6	10.05	16	19.54	21.59	*	*	-	-
	PLACE1005502	9.18	6.48	6.37	3.45	3.87	4.29	4.21	4.81	4.94	*	*	-	-
	PLACE1005513	33.31	27.28	25.19	10.65	9.46	7.45	5.15	5.68	9.33	**	**	-	-
	PLACE1005515	11.7	9.47	10.48	3.66	3.88	4.7	6	4.7	5.35	**	**	-	-
	PLACE1005519	8.43	6.62	7.72	4.49	2.77	2.81	3.1	2.36	5.05	**	*	-	-
	PLACE1005526	8.8	10.4	11.16	2.78	4.21	2.23	3.21	3.51	3.18	**	**	-	-
10	PLACE1005528	26.91	22.55	23.87	12.32	10.17	9.81	19.08	11.09	17.43	**	*	-	-
	PLACE1005530	20.34	15.73	21.3	12.14	14.27	9.82	8.48	8.49	4.49	*	**	-	-
	PLACE1005536	18.12	14.92	18.89	3.17	4.7	3.13	5.93	4.22	4.19	**	**	-	-
	PLACE1005539	42.25	35.31	35.69	15.02	20.49	15.33	32.11	27.06	37.95	**		-	-
	PLACE1005543	35.86	20.69	14.02	11.25	7.4	11.67	9.96	8.84	11.47	*	*	-	-
	PLACE1005544	4.98	5.73	4.38	2.31	2.15	3.58	3.94	2.43	3.09	*	*	-	-
15	PLACE1005550	26.58	26.63	19.78	16.15	16.36	16.32	11.17	10.41	11.67	*	**	-	-
	PLACE1005554	23.49	15.23	16.27	7.02	8.22	4.71	11.04	7.45	10.06	*	*	-	-
	PLACE1005557	10.41	9.97	10.58	3.15	3.37	2.55	4.57	1.9	2.76	**	**	-	-
	PLACE1005563	10.87	11.46	9.72	4.64	6.41	6.88	7.08	5.98	5.9	**	**	-	-
	PLACE1005569	27.85	17.24	16.24	8.47	8.97	6.39	9.6	5.75	9.27	*	*	-	-
20	PLACE1005574	29.21	21.31	10.09	12.87	15.06	14.88	14.12	11.93	13.7			-	-
	PLACE1005584	80.18	66.17	69.3	73.19	89.15	68.42	53.92	40.06	44.96	*		-	-
	PLACE1005590	14.81	9.97	11.59	6.51	9.72	8.47	7.14	6.35	7.74	*		-	-
	PLACE1005595	23.18	18.25	21.16	12.01	10.55	9.94	19.07	11.17	15.91	**		-	-
	PLACE1005601	11.31	7.51	15.39	5.61	6.28	8.37	7.21	9.02	6.01	*		-	-
	PLACE1005603	18.98	21.86	14.52	8.39	9.87	6.77	12.21	11.52	14.3	*		-	-
	PLACE1005604	8.92	8.05	5.89	5	4.08	4.67	3.54	2.41	2.72	*	**	-	-
25	PLACE1005611	33.68	22.2	29.76	11.8	16.07	9.26	20.54	21.42	26.07	*		-	-
	PLACE1005622	30.69	27.64	23.31	11.83	18.8	12.91	22.33	27.88	24.08	*		-	-
	PLACE1005623	9.95	5.8	5.03	2.69	3.21	2.46	4.37	5.18	4.94	*		-	-
	PLACE1005630	8.17	9.56	9.09	4.43	7.2	3.43	4.75	3.89	4.7	*	**	-	-
	PLACE1005639	60.6	66.64	49.38	29.59	36.73	29.78	61.43	55.47	68.03	**		-	-
30	PLACE1005646	17.8	13.84	15.64	9.47	9.48	7.41	6.37	4.37	5.79	**	**	-	-
	PLACE1005647	13.8	11.09	12.51	5.66	5.78	5.71	6.46	4.42	5.24	**	**	-	-
	PLACE1005648	28.95	31.24	20.78	11.15	9.99	10.48	23.48	13.79	10.43	**		-	-
	PLACE1005653	16.03	13.03	11.9	5.52	4.03	5.5	8.65	5.02	7.5	*	*	-	-
	PLACE1005656	15.9	11.93	10.98	7.3	8.49	5.09	13.9	11.98	12.97	*		-	-
	PLACE1005659	10.5	8.34	6.31	5.3	4.01	3.82	4.61	3.03	4.74	*	*	-	-
35	PLACE1005660	12.82	10.45	9.32	4.78	5.95	3.56	5.23	2.26	5.04	**	**	-	-
	PLACE1005664	23.43	27.28	25.73	6.86	20.98	19.65	21.23	13.1	15.7	*		-	-
	PLACE1005666	12.43	11.64	13.67	7.79	10.52	6.99	11.1	12.9	13.47	*		-	-
	PLACE1005669	14.08	13.16	10.98	6.36	8.32	3.2	8.01	5.64	11	*		-	-
	PLACE1005682	22.57	20.73	18.81	7.71	10.55	7.1	12.18	6.07	11.14	**	**	-	-
	PLACE1005698	7.18	6.68	8.02	7.04	5.88	4.02	4.11	5.29	4.18	**		-	-
40	PLACE1005708	17.69	17.02	22.05	11.44	15.51	9.8	12.59	9.54	8.25	*	*	-	-
	PLACE1005725	9.73	11.27	13.97	7.09	8.31	6.27	4.4	6.48	6.4	*	*	-	-
	PLACE1005727	10.04	8.34	11.08	6.79	10.76	10.55	4.59	6.83	6.66	*		-	-
	PLACE1005730	13.44	12.68	11.77	6.36	7.19	8.47	5.04	7.92	8.47	**	**	-	-
	PLACE1005736	5.83	5.5	7.74	3.15	14.38	1.73	4.63	3.09	3.33	*		-	-
	PLACE1005739	7.01	13.06	9.75	7.43	7.8	5.14	2.96	4.16	3.47	*		-	-
45	PLACE1005745	8.04	6.14	6.43	4.26	4.82	5.89	4.13	2.63	4.39	*		-	-
	PLACE1005752	128.36	102.07	139.38	89.6	72.86	89.83	103.85	94.56	103.55	*		-	-
	PLACE1005755	13.46	9.22	14.54	4.95	6.97	3.84	5.58	5.65	3.87	*	*	-	-
	PLACE1005756	10.54	16.55	18.56	10.11	10.69	10.1	6.71	10.57	12.53			-	-
	PLACE1005760	5.35	5.71	8.64	9.33	6.58	7.47	2.74	8.03	2.74			-	-
	PLACE1005763	18.11	17.1	15.35	9.45	11.68	9.73	7.29	9.29	9.82	**	**	-	-
50	PLACE1005768	19.31	21.05	13.08	10.64	11.47	13.73	6.13	9.41	9.52	*		-	-
	PLACE1005771	24.23	23.82	24.14	14.86	21.27	9.53	16.05	19.49	20.2	*		-	-
	PLACE1005783	6.96	4.13	7.26	3.95	3.94	3.49	3.16	3.04	5.15			-	-
	PLACE1005799	43.95	40.56	57.56	28.57	25.86	15.96	30.85	26.7	32.18	*	*	-	-
	PLACE1005802	18.73	15.75	12.71	10.65	14.22	15.85	21.48	13.68	14.38			-	-
	PLACE1005803	14.16	12.88	18.69	6.94	7.06	7.31	4.95	5.48	3.9	**	**	-	-
55	PLACE1005804	16.14	16.06	21.67	10.66	8.62	11.86	4.49	6.03	5.63	*	**	-	-
	PLACE1005813	7.59	9.87	17.86	7.9	9.25	7.77	3.93	4.99	5.43			-	-
	PLACE1005815	17.11	19.5	18.51	7.94	18	6.6	9.57	8.13	8.77	**		-	-

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	PLACE1005101	11.5	9.29	11.94	9.18	4.8	5.47	5.22	3.69	6.78	*	**	-	-
	PLACE1005102	22.17	25.98	27.54	6.07	7.69	6.94	7.15	5.65	7.82	**	**	-	-
5	PLACE1005108	10.15	7.73	6.02	2.99	6.05	3.33	4.8	7.23	6.47				
	PLACE1005110	3.97	2.39	3.45	3.59	3.78	1.51	2.14	1.33	2.15				
	PLACE1005111	23.94	21.69	16.98	14.04	21.5	8.57	20.32	16.74	23.71				
	PLACE1005123	16.94	14.46	12.96	5.45	9.5	5	6.89	4.29	5.82	*	**	-	-
	PLACE1005124	9.14	6.62	6.15	2.06	5.47	1.8	6.61	3.46	5.33	*		-	-
10	PLACE1005128	6.83	4.81	4.87	1.09	3.42	1.91	3.3	3.63	2.82	*	*	-	-
	PLACE1005130	12.43	17.68	12.68	4.58	11.05	5.79	12.55	5.44	10.87				
	PLACE1005141	6.07	3.99	5.18	6.23	2.88	1.1	9.65	2.17	2.82				
	PLACE1005146	10.52	9.7	6.88	6.24	10.69	5.88	5.25	8.14	5.49				
	PLACE1005152	91.15	76.64	75.79	51.06	95.95	52.59	42.4	38.1	38.75	**		-	-
	PLACE1005157	5.11	3.75	6.06	2.82	4.86	2.4	1.9	2.79	1.7	*		-	-
	PLACE1005162	24.79	23.25	27.03	17.82	26.77	20.32	10.52	15.51	11.65	**		-	-
15	PLACE1005170	9.55	7.54	6.99	11.08	19.85	14.4	3.82	8.2	5.94				
	PLACE1005176	5.55	5.17	6.85	2.52	4.13	3.01	2.09	2.54	1.26	*	**	-	-
	PLACE1005181	10.56	18.07	15.09	7.46	9.06	8.01	10.91	12.85	10.11	*		-	-
	PLACE1005184	3.64	4.4	5.5	5.28	11.77	4.26	3.84	3.69	3.06				
	PLACE1005186	5.96	5.07	5.66	2.88	2.54	3.26	3.64	4.33	4.72	**	*	-	-
20	PLACE1005187	20.53	14.71	18.55	9.5	10.66	10.36	12.23	12.87	7.61	*	*	-	-
	PLACE1005189	8.28	7.72	11.92	4.24	7.19	3.99	6.84	7.52	5.99				
	PLACE1005193	39.26	44.11	39.59	31.34	38.66	28.91	36.03	40.56	33.69				
	PLACE1005200	1391.4	1317.2	1202.3	1770.3	2457.3	1744.3	1189.8	1397.9	586.38	*		+	
	PLACE1005205	48.27	41.57	88.17	48.38	52.86	50.82	27.19	13.53	58.32				
	PLACE1005216	10.1	9.74	10.26	5.79	10.41	5.95	3.4	3.26	4.05		**	-	-
25	PLACE1005223	36.86	44.94	37.06	23.02	26.39	16.59	23.41	26.05	19.76	*	**	-	-
	PLACE1005225	5.34	5.81	6.83	4.25	3.02	5.27	2.59	3.79	3.7	*		-	-
	PLACE1005232	37.45	31.95	45.29	20.08	17.68	11.81	24.42	25.92	23.95	**	*	-	-
	PLACE1005239	7.93	6.66	12.45	5.29	3.66	4.36	3.81	3.89	4.86				
	PLACE1005243	16.13	27.58	27.13	19.28	15.8	18.25	6.8	13.48	9.67	*		-	-
	PLACE1005250	5.37	6.75	7.77	5.96	4.93	3.55	3.78	5.49	3.24				
30	PLACE1005261	61.12	61.95	56.67	47.53	36.14	42.92	20.88	23.52	21.14	**	**	-	-
	PLACE1005266	7.03	6.91	9.71	5.82	8.14	5.96	4.18	3.69	5.53	*		-	-
	PLACE1005271	8.47	8.61	6.86	4.77	7.74	3.27	3.65	3.57	3.35	**		-	-
	PLACE1005277	20.92	15.1	22.31	9.09	13.8	9.19	14.68	15.64	18.36	*		-	-
	PLACE1005287	15.1	8.12	14.31	5.25	6.44	6.19	5.01	4.78	3.02	*	*	-	-
	PLACE1005299	10.04	6.09	8.79	4.5	5.75	7.77	4.03	4.98	7.58				
35	PLACE1005305	8.97	7.38	9.77	4.47	3.45	4.18	3.14	3.99	3.81	**	**	-	-
	PLACE1005307	12.01	9.78	33.93	6.5	6.01	6.92	5.78	5.91	8.37				
	PLACE1005308	6.72	9.28	13.18	4.37	5.39	4.05	5.69	4.54	7.02				
	PLACE1005313	8.48	6.04	10.79	5.16	6.95	3.74	4.51	3	4	*		-	-
	PLACE1005320	9.29	12.54	14.31	7.32	9.29	5.14	4.92	4.65	5.56	**		-	-
40	PLACE1005327	31	25.4	35.29	16.09	17.4	11.88	9.82	14.36	14.83	**	**	-	-
	PLACE1005331	13.95	13.36	13.69	9.09	7.58	7.27	7.77	10.5	7.36	**	**	-	-
	PLACE1005335	11.38	7.64	13.28	5.94	7.87	3.65	8.18	25.07	9.78				
	PLACE1005336	29.59	23.98	23.59	14.99	18	12.79	23.63	28.14	27.34	*		-	-
	PLACE1005351	11.96	8.58	19.03	4.6	4.51	3.68	3.27	6.24	4.61	*		-	-
	PLACE1005366	73.8	70.61	70.61	45.9	65.72	42.52	55.33	53.05	50.72	*	**	-	-
	PLACE1005373	9.8	6.9	15.51	3.87	3.73	2.69	2.9	3.58	2.99	*	*	-	-
45	PLACE1005374	31.25	37.89	30.27	12.29	20.74	12.12	33.17	26.39	29.29	**		-	-
	PLACE1005383	9.15	7.91	6.27	5.38	5.79	4.82	3.77	5.77	7.02	*		-	-
	PLACE1005388	14.78	7.34	14.13	9.31	7.11	6.02	6.28	5.49	5.89				
	PLACE1005409	11.86	10.31	14.09	4.95	5.84	5.2	5.09	4.29	5.35	**	**	-	-
	PLACE1005410	13.35	9.74	14.55	5.14	6.18	4.53	6.86	5.79	5.13	**	*	-	-
	PLACE1005426	10.65	4.8	11.31	4.23	3.51	3	2.46	6.49	1.78				
50	PLACE1005431	8.27	7.3	10.2	3.04	3.82	2.32	3.59	3.79	5.31	**	*	-	-
	PLACE1005453	7.65	9.23	18.96	3.09	5.38	3.63	2.3	4.42	3.43				
	PLACE1005467	10.75	11.01	23.28	4.33	7.93	7.46	6.29	7.36	5.29				
	PLACE1005471	7.1	9.43	5.6	3.8	6.04	3.15	3.31	6.18	12.95				
	PLACE1005476	29.52	27.39	29.6	15.57	12.74	15.37	14.04	14.17	16.62	**	**	-	-
	PLACE1005477	9.37	7.91	13.74	3.68	6.31	4.63	5	3.74	6.24	*	*	-	-
55	PLACE1005480	7.1	8	8.36	5.61	4.81	4.45	4.92	2.94	2.27	**	**	-	-
	PLACE1005481	10.21	6.44	10.18	3.66	4.76	4.41	5.76	4.63	3.43	*	*	-	-
	PLACE1005494	9.19	9.94	10.69	3.14	4.05	3.41	4.87	3.69	4.02	**	**	-	-

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	PLACE1004740	5.7	7.45	13.2	8.48	9.7	6.78	5.51	4.01	4.39				
	PLACE1004743	25.2	29.5	32.11	15.32	29.52	15.01	26.22	23.41	28.91				
5	PLACE1004751	26.3	37.3	36.23	17.86	40.34	16.49	27.84	27.38	23.37				
	PLACE1004757	27.21	29.89	35.01	20.07	11.76	13.22	19.63	14.54	16.78	*	**	-	-
	PLACE1004761	36.62	28.57	36.7	16.84	22.51	18.17	11.71	16.47	11.41	**	**	-	-
	PLACE1004773	6.34	3	4.53	3.16	3.36	2.2	2.15	2.55	2.95				
	PLACE1004775	6.98	4.5	11.09	2.47	5.26	2.58	2.74	2.24	1.32				
	PLACE1004777	22.12	29.48	36.12	13.38	18.93	13.39	17.54	20.58	18.98	*		-	-
10	PLACE1004793	38.61	50.6	39.99	29.77	37.64	25.63	24.47	24.63	21.5		**	-	-
	PLACE1004796	8.45	9.48	8.63	3.38	9.79	3.2	3.16	2.68	3.41		**	-	-
	PLACE1004804	5.96	3.66	3.88	1.07	6.07	2.59	2.15	2.35	2.21	*		-	-
	PLACE1004813	7.54	8.6	8.99	4.2	5.66	4.51	4.87	4.35	4.23	**	**	-	-
	PLACE1004814	4.13	4.9	5.89	3.27	11.52	2.25	2.11	31.08	4.24				
	PLACE1004815	5.57	3.05	6.31	3.48	4.55	1.54	4.88	2.03	1.26				
15	PLACE1004816	10.65	12.52	15.46	46.15	13.59	7.88	11.01	6.49	24.95				
	PLACE1004824	10.1	10.18	14.93	3.84	7.8	4.37	6.87	4.52	8.59	*		-	-
	PLACE1004827	6.33	6.31	10.2	1.77	3.73	0.89	1.94	4.36	3.45	*	*	-	-
	PLACE1004836	18.51	24.59	14.94	5.86	11.83	5.72	3.63	5.6	4.83	*	**	-	-
	PLACE1004838	32.68	37.85	26.97	17.25	33.96	16.53	18.4	22.65	17.82	*		-	-
20	PLACE1004840	26.67	33.84	24.99	11.9	21.32	10.63	17.68	19.77	20.11	*	*	-	-
	PLACE1004842	29.36	29.36	36.71	26.03	17.22	20.39	11.25	16.65	13.58	*	**	-	-
	PLACE1004850	12.96	12.93	16.16	9.69	10.5	6.77	8.83	9.34	8.77	*	**	-	-
	PLACE1004868	19.8	16.22	20.8	8.53	16.93	9.03	11.19	11.11	9.47	**	**	-	-
	PLACE1004885	7.93	7.22	6.94	1.5	3.78	3.6	1.94	2.3	1.73	**	**	-	-
	PLACE1004886	8.48	8.07	17.49	4.04	4.63	2.9	4.53	4.47	2.5				
	PLACE1004887	11.79	13.7	11.52	6.38	10.2	7	3.69	4.3	5.52	*	**	-	-
25	PLACE1004896	6.25	6.09	7.72	0.93	8.2	4.59	2.58	2.25	1.01	**	**	-	-
	PLACE1004900	11.28	8.08	8.69	8.58	6.42	4.7	4.96	5.45	3.62	*		-	-
	PLACE1004902	7.13	7.38	10.7	4.87	6.92	2.81	4.29	4.21	2.64	*		-	-
	PLACE1004904	23.08	23.38	37.33	15.52	22.71	12.92	16.1	22.74	25.02				
	PLACE1004911	11.74	10.67	12.36	7.47	12.66	7.84	6.28	5.55	4.09		**	-	-
	PLACE1004913	12.82	12.15	14.37	4.58	7.86	4.67	7.24	7.07	6.68	**	**	-	-
30	PLACE1004918	17.31	18.68	18.94	7.36	10.01	6.44	8.32	5.56	6.18	**	**	-	-
	PLACE1004930	6.79	6.06	5.6	1.3	2	3.65	2.19	1.6	1.33	**	**	-	-
	PLACE1004934	14.99	19.19	12.29	5.42	10.92	8.11	4.61	8.66	7.47	*	*	-	-
	PLACE1004937	18.93	13.85	13.4	11.81	10.98	11.51	15	13.14	16.45				
	PLACE1004949	5	5.22	6.64	5.79	3.31	3.09	4.44	4.05	3.5	*		-	-
35	PLACE1004969	13.91	13.85	11.55	11.71	12.45	9.25	4.82	9.25	5.58	*		-	-
	PLACE1004970	28.75	33.22	29.53	13.51	29.35	23.54	15.11	12.88	14.15	**	**	-	-
	PLACE1004972	13.26	11.27	10.14	5.85	6.98	5.82	4.89	3.42	1.54	**	**	-	-
	PLACE1004974	16.47	29.85	28.89	16.44	23.94	17.53	14.71	17.1	12.72				
	PLACE1004975	12.92	11.66	12.08	5.89	13.8	6.88	5.54	3.55	3.43	**		-	-
	PLACE1004979	7.78	7.47	6.98	3.07	7.65	3.76	4.2	4.56	6.5	*		-	-
40	PLACE1004982	30.58	23.61	26.29	14.3	19.23	11.42	16.82	24.16	30.61	*		-	-
	PLACE1004985	29.68	24.94	25.1	15.66	19.26	12.6	21.78	17.15	20.28	*	*	-	-
	PLACE1005003	16.43	14.5	16.78	6.69	10.88	11.47	1.79	5.8	6.41	*	**	-	-
	PLACE1005004	5.26	4.21	4.06	1.44	3.4	2.25	2.76	1.23	1.09	*	*	-	-
	PLACE1005005	17.79	13.99	18.79	11.99	10.57	8.86	7.49	5.3	5.99	*	**	-	-
	PLACE1005011	10.94	14.04	9.8	3.35	9.24	6.44	3.46	2.32	0.6		**	-	-
45	PLACE1005026	31.74	35.88	28.88	14.82	24.65	14.67	35.74	25.33	28.57	*		-	-
	PLACE1005027	26.29	29.23	15.87	7.97	19.55	9.09	25.84	19.81	24				
	PLACE1005031	38.12	16.05	14.24	9.18	25.55	17.12	7.22	9.47	10.36				
	PLACE1005036	13.86	12.84	17.42	8.68	15.64	10.29	11.23	7.16	6.77	*		-	-
	PLACE1005041	111.13	87.66	72.07	36.83	51	43.47	89.34	93.97	97.87	*		-	-
	PLACE1005046	27.6	29	18.07	15.06	19.65	20.68	22.16	13.25	24				
50	PLACE1005047	50.64	34.98	35.58	25.13	37.56	21.77	26.01	15.91	25.77	*		-	-
	PLACE1005052	10.39	12.55	10.8	4.67	15.46	6.61	10.9	9.05	16				
	PLACE1005055	24.31	33.8	18.61	16.75	10.79	11.1	19.54	11.46	10.17				
	PLACE1005066	7.67	8.41	7	12.21	61.68	20.27	25.44	4.84	8.48				
	PLACE1005077	9.62	6.97	5.62	4.44	4.68	3.2	5.42	5.71	7.8				
	PLACE1005085	10.11	4.46	3.84	3.42	3.7	2.05	2.8	5.01	2.86				
55	PLACE1005086	11.97	11.74	11.67	7.31	18.85	7.84	6.74	6.63	9.62	*		-	-
	PLACE1005088	86.57	83.56	74.65	79.4	89.3	75.33	56.8	52.44	57.53	**		-	-
	PLACE1005089	6.31	4.13	4.27	1.8	3.13	4.07	2.18	0.55	2.04	*		-	-

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	PLACE1004405	6.83	7.82	8.42	1.74	2.87	2.17	1.79	1.77	1.06	**	**	-	-
	PLACE1004407	11.75	11.43	14.16	6.35	10.59	6.28	6.13	6.07	6.47	*	**	-	-
	PLACE1004424	6.94	8.4	8.66	2.02	6.19	2.66	3.43	3.43	2.34	*	**	-	-
5	PLACE1004425	15.08	13.72	13.35	6.65	11.37	7.31	11.7	10.39	11.17	*	*	-	-
	PLACE1004427	9.79	7.13	8.53	5.79	4.82	4.58	2.31	1.96	2.72	*	**	-	-
	PLACE1004428	8.46	5.33	5.57	3.25	3.29	3.21	2.85	2.65	3.1	*	*	-	-
	PLACE1004433	6.57	5.12	7.26	2.15	3.61	2.86	2.76	2.36	3.32	*	**	-	-
	PLACE1004435	5.31	5.07	8.8	2.76	3.41	3.61	2.83	0.69	1.36	*	*	-	-
10	PLACE1004437	123.66	158.84	144.08	116.56	167.95	86.05	49.84	88.42	68.77	**	**	-	-
	PLACE1004441	10.02	12.73	11.55	5.63	11.08	5.24	5.14	4.56	2.63	**	**	-	-
	PLACE1004446	4.95	5.73	4.69	2.19	3.83	1.62	2.79	2.61	2.66	*	**	-	-
	PLACE1004450	17.21	13.77	11.35	7.43	9.82	10.35	13.72	15.81	15.38			-	-
	PLACE1004451	21.93	21.83	22.42	14.24	15.13	14.34	10.39	9.42	10.97	**	**	-	-
	PLACE1004456	5.66	3.89	4.79	2.22	5.14	3.7	4.95	1.66	2.31			-	-
15	PLACE1004458	6.57	4.16	7.77	2.1	5.75	2.16	2.41	3.03	1.54	*	*	-	-
	PLACE1004460	6.93	5.13	4.79	2.01	4.85	1.65	3.61	2.54	1.12	*	*	-	-
	PLACE1004467	10.11	11.82	9.53	2.74	4.49	5.1	3.12	1.44	1.55	**	**	-	-
	PLACE1004471	14.09	22.14	14.79	8.38	18.33	11.21	5.81	8.78	5.83	*	*	-	-
	PLACE1004473	8.27	14.31	9.86	3.42	8.92	3.77	4.46	5.47	8.39			-	-
	PLACE1004475	4.2	2.67	2.47	2.6	8.42	6.68	3.42	2.58	2.7			-	-
20	PLACE1004482	42.46	39.9	32.17	18.98	31.24	19.51	41.1	31.66	45.4	*	*	-	-
	PLACE1004491	3.4	5.37	4.92	2.26	2.87	1.21	3.11	1.77	1.09	*	*	-	-
	PLACE1004492	14.75	10.5	4.26	8.39	16.03	7.27	3.28	2.19	2.31			-	-
	PLACE1004506	18.11	19.82	21.23	12.46	23.49	16.62	17.33	12.61	13.21	*	*	-	-
	PLACE1004507	14.62	14.01	19.19	12.21	20.75	13.12	5.25	3.93	6.09	**	**	-	-
25	PLACE1004510	8.27	7.11	7.77	4.2	4.81	1.36	5.06	2.94	3.49	*	**	-	-
	PLACE1004516	5.75	6.67	3.83	4.51	8.48	2.72	4.69	2.56	3.06			-	-
	PLACE1004518	9.64	6.39	6.6	5.58	7.29	2.45	2.45	4.03	4.48	*	*	-	-
	PLACE1004519	45.05	38.06	29.43	16.22	21.37	17.38	37.02	29.03	42.7	*	*	-	-
	PLACE1004520	5.65	5.88	4.83	5	23.3	3.26	7.89	2.48	5.32			-	-
	PLACE1004530	8.55	8.01	8.19	4.33	4.87	3.07	5.74	4.33	5.72	**	**	-	-
30	PLACE1004545	18.35	9.42	10.89	5.28	6.16	4.82	7.69	5.29	6.88	*	*	-	-
	PLACE1004547	8.88	4.99	7.28	3.79	10.69	2.44	2.07	2.03	2.03	*	*	-	-
	PLACE1004548	8.06	7.34	7.17	1.99	3.55	1.53	3.36	1.74	4.09	**	**	-	-
	PLACE1004550	9.29	4.74	4.57	6.39	7.24	2.64	11.12	3.11	5.12			-	-
	PLACE1004551	28.29	24.92	17.22	13.36	20.97	8.68	22.61	16.23	21.2			-	-
	PLACE1004559	38.64	25.35	19.12	16.44	15.14	16.5	25.43	24.95	33.62			-	-
35	PLACE1004562	50.29	46.32	50.21	54.16	56.41	39.84	35.71	42.24	45.43			-	-
	PLACE1004564	12.01	9.86	9.29	4.62	9.04	5.79	9.27	6.54	10.21			-	-
	PLACE1004604	17.66	17.86	15.11	9.74	9.81	7.66	16.75	11.47	16.51	**	**	-	-
	PLACE1004611	6.51	3.64	4.04	1.34	7.9	1.38	3.77	0.9	3.89			-	-
	PLACE1004629	7.22	4.23	7.21	2.88	5.64	1.27	4	2.37	6.88			-	-
40	PLACE1004630	5.71	5.71	5.05	2.41	3.38	0.92	6.59	3.07	5.19	*	*	-	-
	PLACE1004637	15.22	15.19	15.73	4.5	6.73	18.68	5.83	11.18	5.64	*	*	-	-
	PLACE1004645	16.19	10.08	17.1	10.36	11.57	13.97	10.9	12.36	6.68			-	-
	PLACE1004646	4.97	5.64	4.39	4.13	5.71	3.39	2.58	3.34	2.67	**	**	-	-
	PLACE1004648	5.49	4.11	3.96	3.64	5.16	4.29	1.91	1.5	0.14	**	**	-	-
	PLACE1004655	3.31	1.36	2.13	1.15	3.39	0.72	0.74	1.06	1.47			-	-
	PLACE1004658	61.58	67.15	67	38.54	49.97	38.25	52.62	54.12	65.08	**	**	-	-
45	PLACE1004664	1.19	4.36	2.93	2.86	5.73	2.7	1.85	2.13	1.51			-	-
	PLACE1004672	28.88	30.42	25.6	16.98	20.16	13.59	16.28	11.87	13.82	**	**	-	-
	PLACE1004674	4.27	1.6	4.11	2.41	3.15	1.89	1.56	2.96	3.87			-	-
	PLACE1004681	6.54	6.41	13.08	7.27	21.72	12.91	4.64	4.9	3.36			-	-
	PLACE1004686	4.34	3.3	4.61	2.4	3.38	2.38	2.29	3.63	0.39			-	-
	PLACE1004690	6.4	6.12	8.6	4.32	13.96	8.66	2.73	4.56	2.78	*	*	-	-
50	PLACE1004691	12.44	14.52	17.78	7.94	13.42	7.13	8.04	12.14	11.56			-	-
	PLACE1004693	8.4	10.27	9.88	5.82	5.16	5.28	2.45	3.25	2.73	**	**	-	-
	PLACE1004701	144.88	198.16	97.68	190.88	167.91	84.32	88.33	73.95	139.9			-	-
	PLACE1004705	3.47	3.01	2.93	2.39	3.96	1.27	1.92	3.49	3.87			-	-
	PLACE1004708	48.88	43.13	45.03	31.98	28.55	23.56	15.06	39.44	11.77	**	**	-	-
	PLACE1004716	20.04	14.83	24.39	8.19	12.11	8.27	8.37	8.78	9.57	*	*	-	-
55	PLACE1004722	5.57	7.39	10	4.45	6.54	3.93	3.57	3.69	3.48	*	*	-	-
	PLACE1004736	16.71	17.15	24.67	12.34	13.91	6.89	9.19	22.16	13.53			-	-
	PLACE1004737	5.22	7.74	7.78	3.83	6.24	1.88	3.06	3.38	3.05	*	*	-	-

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	PLACE1003900	9.05	5.59	5.53	3.41	9.33	4.49	4.41	6.87	4.79				
	PLACE1003902	9.43	6.06	7.46	2.35	3.67	1.76	3.96	1.76	3.05	*	*	-	-
5	PLACE1003903	17.33	19.36	14.55	9.42	10.87	10.03	9.08	6.56	7.77	**	**	-	-
	PLACE1003915	14.67	18	11.19	12.64	10.36	9.98	8.5	5.29	8.8		*	-	-
	PLACE1003918	10.58	6.62	8.73	7.34	5.29	3.45	7.05	5.69	4.94			-	-
	PLACE1003923	11.24	8.95	4.47	2.95	4.6	4.65	4.65	3.26	3.87			-	-
	PLACE1003932	9.02	5.88	4.39	2.18	9.39	4.8	6.3	4.31	5.14			-	-
	PLACE1003936	11.92	7.46	8.85	2.47	4.23	4.36	9.51	9.11	10.65	*		-	-
10	PLACE1003966	11.6	10.3	9.01	5.2	6.1	4.37	6.77	5.83	6.22	**	**	-	-
	PLACE1003968	4.88	5.07	4.71	0.66	3.28	0.8	3.51	1.87	0.72	*	*	-	-
	PLACE1004018	4.89	2.45	2.55	4.7	8.1	2.19	2.99	1.85	1.89			-	-
	PLACE1004020	12.24	14.87	17.86	7.54	8.19	5.65	7.35	9.99	10.06	*	*	-	-
	PLACE1004028	2.49	1.88	2.35	1.78	7.14	2.28	1.01	1.49	0.82	*		-	-
	PLACE1004034	6.5	4.91	7.64	2.7	5.45	2.86	3.71	3.07	1.75	*		-	-
15	PLACE1004042	5.84	2.98	4.53	2.04	4.08	3.55	3.58	3.9	2.95			-	-
	PLACE1004078	6.26	4	5.13	2.35	3.57	1.92	1.94	3.66	4.31	*		-	-
	PLACE1004103	4.16	6.73	8.68	2.05	3.45	2.36	1.89	0.91	3.01	*	*	-	-
	PLACE1004104	0.6	1.12	3	1.89	7.2	1.01	1.88	2.79	1.52			-	-
	PLACE1004113	0.85	1	0.69	0.62	4.18	1	1.54	0.77	1.59			-	-
20	PLACE1004114	12.27	11.63	12.84	9.04	6.13	6.15	17.17	8.3	11.59	**		-	-
	PLACE1004118	25.56	29.6	34.14	10.69	22.13	12.84	16.2	19.15	19.4	*	*	-	-
	PLACE1004128	6.77	8.44	8.16	3.47	7.23	7.78	2.81	3.08	3.14		**	-	-
	PLACE1004130	9.36	4.88	10.96	8.2	7.53	9.87	4.28	6.71	6.1			-	-
	PLACE1004149	2.51	2.43	6.21	2.78	5.35	2.6	3.3	3.26	2.01			-	-
	PLACE1004156	13.06	9.08	13.29	7.39	11.05	6.74	9.05	9.4	10.62			-	-
25	PLACE1004160	20.97	19.94	23.27	14.03	30.3	7.33	16.97	16.65	14.66	*		-	-
	PLACE1004161	7.06	8.18	8.06	5.83	7.98	4.77	4.77	4.44	4.99	**		-	-
	PLACE1004166	41.1	39.11	42.06	19.36	22.19	21.87	24.56	28.32	30.02	**	**	-	-
	PLACE1004168	32.92	30.07	31.55	16.2	15.66	13.35	21.23	25.32	27.11	**	*	-	-
	PLACE1004170	14.53	21.92	19.94	10.32	12.85	10.61	11.03	18.31	8.2	*		-	-
	PLACE1004176	2.75	3.62	3.97	2.55	6.19	3.35	0.34	1.55	0.72		**	-	-
	PLACE1004183	36.95	43.53	45.99	27.78	51.84	25.82	21.25	25.43	23.92	**		-	-
30	PLACE1004197	5.75	5.24	7.84	2.14	4.97	3.74	3.04	2.49	3.28	*		-	-
	PLACE1004199	21.75	27.66	12.12	12.42	24.13	11.64	10.47	10.88	12.46			-	-
	PLACE1004203	11	13.62	10.81	4.23	10.81	6.59	10.74	11.32	12.32			-	-
	PLACE1004242	20.49	16.48	21.11	6.81	6.93	13.71	6.23	11.27	9.24	*	**	-	-
	PLACE1004249	6.73	11	11.25	4.47	10.26	6.2	5.79	8.22	6.09			-	-
35	PLACE1004255	32.98	36.35	38.9	17.52	26.48	21.84	27.95	27.86	29.02	*	*	-	-
	PLACE1004256	12.54	10.61	15.03	5.99	5.51	6.23	3.84	5.04	3.23	**	**	-	-
	PLACE1004257	3.25	14.27	11.31	3.36	8.12	8.15	2.7	9.49	4.03			-	-
	PLACE1004258	11.88	26.37	21.62	8.76	18.18	12.25	12.69	13.44	9.66			-	-
	PLACE1004270	8.17	8.34	11.88	6.47	6.77	11.76	8.27	3.74	6.39			-	-
	PLACE1004272	1.52	1.56	3.17	1.04	3.51	0.46	1.56	1.06	1.78			-	-
40	PLACE1004273	23.31	38.14	20.87	17.38	14.34	7.01	17.2	17.07	20.56			-	-
	PLACE1004274	12.13	13.19	15.06	6.12	12.19	6.04	6.47	11.78	6.75			-	-
	PLACE1004277	18.65	17.3	16.15	12.2	21.29	13	13.46	15.12	20.1			-	-
	PLACE1004279	20.99	17.87	19.27	8.96	30.05	13	8.21	11.3	11.19		**	-	-
	PLACE1004282	12.65	12.53	16.59	4.95	9.12	5.25	6.7	6.34	5.89	*	**	-	-
	PLACE1004284	9.58	14.66	15.53	8.85	7.35	5.65	6.2	6.82	3.1	*	*	-	-
45	PLACE1004289	11.29	10.79	11.74	3.16	5.71	3.35	2.35	6.14	2.14	**	**	-	-
	PLACE1004298	19.52	18.15	16.36	8.87	22.44	9.43	11.73	9.62	8.57		**	-	-
	PLACE1004302	6.5	4.57	6.26	4.18	7	3.55	1.56	4.16	0.55	*		-	-
	PLACE1004305	7.16	4.88	7.78	4.02	11.98	4.38	3.44	4.62	2.94	*		-	-
	PLACE1004316	21.08	20.32	25.43	14.83	15.07	12.7	13.59	13.17	12.8	*	**	-	-
	PLACE1004322	9.54	3.44	10.02	13.05	22.34	1.36	2.52	7.35	3.43			-	-
50	PLACE1004325	13.79	15.32	21.15	8.25	8.41	6.63	10.63	11.11	9.78	*		-	-
	PLACE1004332	13.76	26.81	27.83	10.52	16.97	4.86	9.28	8.37	7.2	*		-	-
	PLACE1004336	9.25	9.04	10.31	3.17	5.77	3.05	2.53	4.29	2.93	**	**	-	-
	PLACE1004346	11.14	15.86	16.19	8.6	6.7	8.17	14.67	2.81	3.65	*		-	-
	PLACE1004358	4.9	6.6	5.3	4.66	3.48	2.52	2.87	2.07	1.85		**	-	-
	PLACE1004376	4.37	7.66	6.28	3.82	16.23	3.24	18.46	3.42	3.06			-	-
55	PLACE1004384	20.82	15.88	20.1	10.2	14.54	11.76	10.47	11.55	9.77	*	**	-	-
	PLACE1004385	18.21	25.32	29.3	6.24	11.85	7.12	18.49	9.57	9.67	*		-	-
	PLACE1004388	13.03	15.64	14.98	6.25	11.49	8.38	7.85	6.97	4.61	*	**	-	-

Table 474

	PLACE1003549	16.21	19.52	27.74	11.72	18.71	13.09	12.65	15.02	13.74			
	PLACE1003553	12.63	20.69	18.89	10.17	18.28	8.51	8.12	9.44	8.9	*	-	
	PLACE1003566	9.49	12.35	9.77	7.53	10.49	7.08	3.77	4.66	6.9	*	-	
5	PLACE1003568	5.3	4.9	19.78	3.8	7.59	3.87	5.3	6.3	4.21			
	PLACE1003573	32	27.25	37.46	17.52	23.28	16.23	15	18.28	19.13	*	**	-
	PLACE1003575	66.07	60.03	70.99	42.8	62.9	38.6	75.17	59.46	62.58			
	PLACE1003583	8.33	4.65	6.88	3.47	6.26	5.07	2.38	3.4	4.1	*	-	
	PLACE1003584	12.09	10.66	18.02	7.17	8.48	19.2	8.54	7.15	2.79			
10	PLACE1003592	29.77	33.55	28.89	21.34	26.31	15.85	12.59	8.49	12.12	*	**	-
	PLACE1003593	7.15	3.54	8.27	3.34	2.24	1.83	3.11	5.19	5.45			
	PLACE1003594	7.18	8.49	9.31	5.03	6.61	3.38	4.99	3.18	2.37	*	**	-
	PLACE1003596	5.11	3.13	4.25	2.82	5	3.14	3.75	4.1	3.76			
	PLACE1003598	31.57	29.75	31.55	14.81	20.6	17.31	5.31	17.81	7.34	**	**	-
	PLACE1003602	6.51	2.47	7.47	1.84	4.91	0.98	2.4	4.6	2.56			
15	PLACE1003605	10	6.14	7.29	5.28	8.85	3.75	6.45	5.07	2.93			
	PLACE1003611	15.94	16.54	24.25	11	9.1	12.46	10.14	10.62	12.83	*	-	
	PLACE1003618	7.78	7.74	12.1	3.49	6.12	2.68	4.58	5.29	6.93	*	-	
	PLACE1003625	14.19	21.53	18.27	11.53	22.81	17.82	13.8	9.54	11.86			
	PLACE1003626	11.55	14.38	14.6	5.94	11.97	7.26	4.57	4.49	2.04	**	-	
	PLACE1003630	3.88	4.65	3	4.62	4.58	3.02	3.68	3.68	1.66			
20	PLACE1003635	8.33	7.29	10.19	5.35	5.59	4.6	3.51	4.48	4.17	*	**	-
	PLACE1003638	4.33	3.09	4.93	3.86	6.02	2.38	2.01	3.6	1.35			
	PLACE1003644	35.03	24.77	24.25	19.14	20.11	17.95	8.77	10.6	6.97	**	-	
	PLACE1003654	75.56	74.57	74.52	77.61	72.17	67	35.39	33.92	31.06	**	-	
	PLACE1003656	7.81	5.19	8.37	3.06	4.32	3.8	1.16	1.63	2.96	*	**	-
	PLACE1003680	16.25	21.8	21.11	12.32	17.1	13.06	14.44	12.72	12.99	*	-	
25	PLACE1003689	12.05	8.79	12.65	4.93	9.33	7.48	3.9	4.69	4.41	**	-	
	PLACE1003670	110.15	83.05	76.99	55.55	52.05	50.36	48.6	63.39	52.6	*	*	-
	PLACE1003671	11.47	3.49	4.69	7.13	4.59	4.35	3.71	2.69	2.22			
	PLACE1003697	9.65	10.83	16.05	9.21	10.08	8.25	6.51	4.97	6.76	*	-	
	PLACE1003704	10.59	7.92	10.14	6.15	6.81	6.51	4.22	3.63	3.86	*	**	-
	PLACE1003709	9.06	8.99	10.78	5.34	6.7	5.47	3.53	3.36	1.83	**	**	-
30	PLACE1003711	28.05	31.28	32.33	21.12	31.77	27.07	16.12	16.73	14.21	**	-	
	PLACE1003723	27.32	34.97	28.91	18.16	26.8	21.24	17.63	17.4	10.81	**	-	
	PLACE1003724	29.33	36.12	25.65	19.94	34.26	21.71	26.66	13.83	15.71			
	PLACE1003737	8.24	6.31	9.31	2.89	4.06	2.72	3.31	2.56	3.28	**	**	-
	PLACE1003738	5.02	4.71	5.03	8.78	2.78	9.43	1.8	2.2	2.93	**	**	-
	PLACE1003742	17.85	18.69	20.82	14.24	7.57	8.14	10.49	8.98	8.16	*	**	-
35	PLACE1003744	5.48	2.82	3.4	0.66	1.04	1.08	1.74	1.94	0.38	*	-	
	PLACE1003758	23.32	22.73	21.79	9.9	15.26	16.38	15.29	14.75	13.36	*	**	-
	PLACE1003760	40.1	57.96	44.54	28.85	37.15	22.14	11.09	14.54	11.41	**	-	
	PLACE1003762	65.35	58.09	77.98	35.3	88.27	46.92	46.48	32.75	38.3	*	-	
	PLACE1003765	93.01	107	70.69	55.75	97.14	46.35	33.33	44.53	44.51	*	-	
40	PLACE1003768	34.07	21.23	23.34	10.87	13.6	6.67	25.2	17.43	28.3	*	-	
	PLACE1003771	5.38	5.57	5.27	3.47	7.27	2.71	1.77	1.77	2.44	**	-	
	PLACE1003772	5.63	7.96	4.89	1.17	7.74	2.12	3.12	2.24	2.28	*	-	
	PLACE1003783	5.15	3.84	3.95	2.36	3.83	1.92	1.65	1.17	0.84	**	-	
	PLACE1003784	5.59	6.96	5.68	2.81	4.01	2.73	3.1	2.42	1.88	**	**	-
	PLACE1003788	34.98	33.59	27.31	19.49	15.1	21.27	12.2	5.92	10.73	*	**	-
45	PLACE1003795	26.48	33.02	23.72	12.5	28.84	12.28	23.33	16.94	16.45			
	PLACE1003827	9.74	8.51	10.55	21.41	16.48	10.33	5.7	3.14	1.91	**	-	
	PLACE1003833	14.14	10.78	12.27	5.74	8.62	5.66	5.59	6.29	7.37	*	**	-
	PLACE1003839	259.31	241.18	222.89	217.52	173.45	205.17	163.55	139.4	151.61	**	-	
	PLACE1003845	69.72	60.2	118.04	40.89	28.14	26.89	48.08	43.86	45.73			
	PLACE1003850	9.53	11.45	9.42	3.3	8.43	5.88	6.51	3.87	5.97	*	-	
50	PLACE1003852	32.49	28.32	26.73	21.76	30.69	26.63	18.45	14.12	17.92	**	-	
	PLACE1003858	19.9	13.83	13.86	8.03	12.82	5.24	7.08	4.19	7.71	*	-	
	PLACE1003861	40.22	66.3	33.51	12.47	57.37	18.72	33.23	18.67	19.13			
	PLACE1003864	34.51	84.76	122.59	61.83	70.16	65.03	69.32	56.31	42.37			
	PLACE1003870	9.43	6.5	5.9	2.94	3.32	3.4	5.69	3.86	5.74	*	-	
	PLACE1003885	10.07	6.76	3.91	3.5	6.22	3.91	3.49	3.72	6.33			
55	PLACE1003886	12.66	12.76	10.57	6.28	9.54	3.51	13.66	8.88	12.28	*	-	
	PLACE1003888	15.48	13.6	17.45	7.94	12.36	6.23	14.56	8.73	9.07	*	-	
	PLACE1003892	6.59	4.85	8.21	5.64	7.14	2.75	3.77	2.15	4.24			

Table 473

	PLACE1003145	12.65	9.94	12.43	3.74	5.65	4.53	6.94	8.05	4.35	**	*	-	-
	PLACE1003147	26.48	26.41	26.84	11.9	16.38	12.11	16.22	18.19	14.71	**	**	-	-
5	PLACE1003153	10.51	10.55	11.78	2.97	6.22	4.35	5.52	7.01	3.61	**	**	-	-
	PLACE1003163	8.08	7.83	4.23	1.17	6.16	2.64	1.92	2.39	2.29	*		-	-
	PLACE1003172	11.82	10	15.87	-6.77	13.23	9.45	7.97	8.79	7.46			-	-
	PLACE1003174	15.37	9.05	12.28	9.12	4.76	2.9	4.07	4.06	4.97	*		-	-
	PLACE1003176	5.18	3.89	6.16	2.81	3.31	2.68	2.96	3.04	2.06	*	*	-	-
	PLACE1003181	9.36	6.29	8.52	1.78	2.48	3.42	2.28	1.97	1.19	**	**	-	-
10	PLACE1003184	23.08	18.89	20.23	9.3	11.84	12.17	13.78	13.13	10.63	**	**	-	-
	PLACE1003190	8.27	8.79	9.4	3.25	4.54	3.59	4.27	2.82	4.68	**	**	-	-
	PLACE1003200	9.79	7.56	9.39	2.41	6.08	2.83	2.42	1.72	3.3	*	**	-	-
	PLACE1003205	126.72	130.26	137.95	113.01	122.92	109.79	99.16	86	89.92	*	**	-	-
	PLACE1003209	2.16	16.18	2.48	0.47	10.18	0.97	2.71	3.36	3.16			-	-
	PLACE1003214	22.58	19.76	20.17	11.54	15.74	15.47	11.96	12.4	14.5	*	**	-	-
15	PLACE1003229	11.83	11.81	10.65	4.92	4.84	3.94	7.34	5.53	5.38	**	**	-	-
	PLACE1003238	8.67	6.17	5.07	3.06	1.84	4.56	1.99	2.39	2.01	*	**	-	-
	PLACE1003249	64.47	55.31	48	33.64	42.61	31.3	33.47	28.86	18.94	*	*	-	-
	PLACE1003256	18.3	25.49	17.26	10.97	13.36	18.51	17.3	11.04	4.08			-	-
	PLACE1003258	49.07	49.35	44.47	34.46	50.32	29.75	23.51	18.19	23.43	**		-	-
	PLACE1003279	10.3	14.58	7.71	5.42	10.68	4.9	6.28	3.11	11.78			-	-
20	PLACE1003294	21.78	16.35	23.18	6.69	11.5	6.25	12.62	8.7	9.35	*	*	-	-
	PLACE1003296	42.17	40.86	35.97	38.78	41.16	32.25	23.79	17.48	17.34	**	**	-	-
	PLACE1003297	49.69	50.94	40.2	15.98	31.29	21.96	19.68	11.27	24.02	*	**	-	-
	PLACE1003302	25.32	31.25	29.06	9.93	15.98	12.55	24.23	16.15	20.93	**	*	-	-
	PLACE1003334	185.2	152.92	141.49	146.89	154.17	125.41	110.94	69.78	76.19	*		-	-
	PLACE1003337	48.98	43.27	41.6	31.2	33.91	22.01	17.95	11.14	16.35	*	**	-	-
25	PLACE1003342	33.19	37.72	34.03	17.42	45.69	20.1	23.58	17.03	23.28	**		-	-
	PLACE1003343	6.84	5.22	10.11	7.16	4.99	3.86	5.46	2.81	3.26			-	-
	PLACE1003344	11.02	7.76	2.99	1.99	8.29	3.43	7.57	2.34	3.75			-	-
	PLACE1003353	19.54	21.69	17.69	8.25	8.74	5.2	10.76	10.8	13.5	**	**	-	-
	PLACE1003361	44.83	37.64	33.66	33.83	37.42	32.13	26.59	20.73	23.72	*		-	-
	PLACE1003366	12	18.42	15.82	7.6	12.12	4.37	10.36	5.92	5.45	*		-	-
30	PLACE1003369	4.81	3.43	3.93	0.94	3.73	1.31	1.63	1.07	0.84	**		-	-
	PLACE1003372	13.64	7.8	10.54	3.88	10.47	3.36	4.52	1.77	3.52	*		-	-
	PLACE1003373	8.88	8.18	9.38	3.94	3.76	0.99	4.29	2.31	3.38	**	**	-	-
	PLACE1003375	22.91	23.46	16.65	11.55	17.67	11.28	29.27	13.77	19.27			-	-
	PLACE1003378	1.92	0.43	0.47	1.45	0.86	1.22	0.77	0.97	0.76			-	-
	PLACE1003383	15.75	12.58	10.55	7.3	8.49	3.81	2.43	7.46	2.15	*	*	-	-
35	PLACE1003394	19.48	13.29	10.26	6.3	12.66	5.09	11.98	10.6	17.14			-	-
	PLACE1003401	8.87	6	5.04	3.66	4.75	2.93	5.66	3.3	5.47			-	-
	PLACE1003405	7.6	8.07	5.91	3.39	5.27	4.58	4.12	1.47	2.3	*	*	-	-
	PLACE1003407	10.24	7.49	7.79	2.43	3.28	2.12	4.16	2.37	3.5	**	**	-	-
	PLACE1003420	7.71	5.57	6.39	3.78	6.36	2.21	2.89	1.88	2.26	**		-	-
	PLACE1003428	6.55	5.9	4.45	2.19	4.16	0.9	3.49	2.31	8.87	*		-	-
40	PLACE1003432	51.6	69.55	75.77	36.76	44.92	44.22	42.27	64.14	52.55	*		-	-
	PLACE1003438	1.98	2.35	1.48	2.21	4.73	1.43	4.08	0.93	0.7			-	-
	PLACE1003452	3.01	3.41	4.3	1.8	3.84	2.08	2.34	2.25	1.64	*		-	-
	PLACE1003454	3.87	4.87	5.58	2.54	4.5	3.92	2.57	2.43	1.33	*		-	-
	PLACE1003455	2.92	2.41	4.04	1.6	1.9	1.64	1.75	0.8	1.88	*	*	-	-
45	PLACE1003456	2.36	4.17	3.63	2.06	10.37	2.22	2.4	1.94	0.64			-	-
	PLACE1003460	4.76	5.09	4.73	4.08	6	2.64	6.38	3.46	2.92			-	-
	PLACE1003478	6.15	2.11	2.66	1.92	3.96	2	5.39	1.91	4.04			-	-
	PLACE1003484	6.84	8.17	8.98	4.36	6.73	3.1	4.12	4.68	4.63	**		-	-
	PLACE1003493	83.14	41.9	88.74	60.01	72.45	45.07	64.55	71.53	53.96			-	-
	PLACE1003503	7.36	5.03	5.29	1.12	3.91	3.51	2.36	3.64	2.4	*		-	-
50	PLACE1003505	32.2	45.22	38.51	20.24	36.17	23.66	24.44	34.5	27.01			-	-
	PLACE1003516	7.02	7.8	7.71	2.88	5.85	3.76	2.24	2.98	3.86	*	**	-	-
	PLACE1003519	4.06	1.81	5.55	3.83	3.85	1.91	2.58	1.64	0.34			-	-
	PLACE1003520	12.72	16.96	9.15	8.71	16.58	8.45	4.85	6.39	5.22			-	-
	PLACE1003521	14.95	10.13	14.17	6.02	7.53	5.56	6.77	8.38	4.75	*	*	-	-
	PLACE1003525	20.17	19.13	25.59	15.85	14.55	12.63	20.97	18.71	22.36	*		-	-
55	PLACE1003528	35.65	25.12	43.38	21.63	30.38	28.64	16.02	24.64	18.65			-	-
	PLACE1003529	11.19	16.69	21.6	20.8	16.66	31.76	18.24	15.7	15.44			-	-
	PLACE1003537	3.15	4.02	8.21	3.24	3.64	3.89	1.76	2.65	2.32			-	-

Table 472

	PLACE1002604	8.27	3.87	5.87	4.56	4.62	4.03	4.08	2.97	4.32			
	PLACE1002612	7.74	6.07	5.48	4.79	5.04	4.26	2.87	1.84	4.68	*		-
	PLACE1002625	8.98	7.38	5.26	3.91	6.23	4.29	4.63	2.42	5.37			
5	PLACE1002638	10.71	6.82	9.31	4.39	4.89	3.67	4.86	2.91	4.12	*	*	-
	PLACE1002655	20.05	17.22	10.12	8.15	9.62	8.61	4.45	4.4	4.69	*		-
	PLACE1002665	2.85	2.11	1.16	1.34	3.18	0.77	1.36	1.74	1.47			
	PLACE1002685	40.11	36.95	37.09	21.27	32.67	18.25	35	28.12	33.61	*		-
	PLACE1002692	10.4	8.99	7.49	3.22	6.32	3.77	7.61	3.76	6.83	*		-
10	PLACE1002714	16.77	11.38	9.27	6.42	9.93	4.45	11.48	11.6	13.87			
	PLACE1002721	75.97	53.95	40.28	41.2	57.12	33.39	36.52	34.39	33.55			
	PLACE1002722	22.13	17.93	14.21	5.69	15.15	8.22	22.92	15.1	22.19			
	PLACE1002726	7.85	7.95	6.76	3.03	4.47	1.55	5.1	2.92	3.72	**	**	-
	PLACE1002756	11.64	12.77	9.75	6.41	12.37	7.42	9.89	7.27	9.98			
	PLACE1002768	14.14	13.66	17.41	6.96	8.91	8.35	8.23	12.96	13.43	**		-
15	PLACE1002772	11.58	12.05	13.28	5.34	9.18	5.42	7.04	6.7	6.74	*	**	-
	PLACE1002775	3.87	3.22	3.82	4.05	3.19	0.98	1.6	5	2.48			
	PLACE1002780	8.73	8.1	8.81	2.91	6.16	2.82	3.15	6.37	4.25	*	*	-
	PLACE1002782	15.09	16.8	18.22	6.13	9.37	3.95	11.73	13.03	12.96	**	*	-
	PLACE1002794	41.39	43.8	46.4	31.76	35.79	20.44	25	22.44	21	*	**	-
	PLACE1002795	26.08	25.22	25.57	11.54	12.89	13.19	17.32	17.1	14.49	**	**	-
20	PLACE1002811	45.78	45.63	45.46	25.72	32.56	23.15	17.78	22.39	22.65	**	**	-
	PLACE1002815	15.82	13.42	19.22	7.58	7.46	5.69	11.61	7.34	8.29	**	*	-
	PLACE1002816	4.86	2.68	5.34	8.68	10.83	8.75	3.44	3.56	2.71	**	+	-
	PLACE1002822	12.2	11.93	11.07	6.15	11.75	10.16	7.41	12.24	8.12			
	PLACE1002833	9.14	11.66	12.33	5.57	7.46	7.07	4.75	8.35	4.08	*	*	-
	PLACE1002834	6.07	5.81	8.84	3.09	3.53	4.27	1.8	2.98	2.41	*	*	-
25	PLACE1002835	58.22	46.91	63.08	49.95	62.63	44.56	42.91	35.34	35.49	*		-
	PLACE1002839	20.38	22.46	21.91	8.31	11	7.79	15.07	14.42	13.25	**	**	-
	PLACE1002851	24.75	15.11	15.09	7.91	14.09	8.63	20.69	22.56	25.92			
	PLACE1002853	17.64	15.34	22.8	5.52	6.45	6.1	6.28	8.18	4.17	**	**	-
	PLACE1002881	24.86	14.92	26.75	13.38	16.28	13.04	14.2	20.46	17.04			
	PLACE1002901	14.25	14.06	17.74	6.8	8.96	8.66	8.78	4.53	4.12	**	**	-
30	PLACE1002904	18.88	26.12	22.33	14.53	22.41	13.83	9.21	13.95	10.54	*		-
	PLACE1002905	4.41	8.93	12	2.54	8.87	4.07	3.83	5.83	4.18			
	PLACE1002908	13.22	16.1	15.63	15.4	12.89	11.14	8.47	5.63	8.35	**		-
	PLACE1002911	10.97	9.2	9.02	6.7	11.96	7.89	5.04	4.71	5.5	**		-
	PLACE1002941	15.88	10.4	17.73	6.14	8.22	5.77	14.58	18	12.89	*		-
	PLACE1002950	23.65	20.5	17.79	16.96	17.44	5.49	24.31	14.61	17.93			
35	PLACE1002955	5	5.58	7.97	3.29	6.71	3.61	3.06	3.96	2.19	*		-
	PLACE1002958	88.1	92.98	103.56	90.74	91.81	78.41	59.31	53.18	47.63	**		-
	PLACE1002962	3.13	4.33	9.15	0.9	4.01	2.47	0.8	0.76	1.87			
	PLACE1002967	35.4	50.14	51.41	21.55	34.48	29.97	31.98	26.21	37.19			
	PLACE1002968	4.93	4.01	6.65	3.18	3.05	1.93	2.74	2.46	2.95	*	*	-
	PLACE1002976	59.32	76.52	110.97	52.57	103.04	77.15	42.49	35.2	48.8			
40	PLACE1002991	6.98	4.41	7.57	2.11	5.56	3.18	2.77	5.13	6.66			
	PLACE1002993	13.31	16.98	19.44	6.54	17.73	10.56	9.53	20.73	7.89			
	PLACE1002996	33.54	26.83	25.33	24.84	36.88	25.62	15.5	28.74	20.58			
	PLACE1003010	5.97	5.37	8.7	4.68	3.28	4.88	3.8	4.41	3.12			
	PLACE1003025	19.13	20.65	29.6	10.42	22.81	11.62	14.02	22.04	12.57			
	PLACE1003027	4.81	4.57	8.94	2.3	4.47	0.55	3.14	5.14	2.31			
45	PLACE1003044	13.45	13	13.59	5.33	7.16	6.74	3.17	4.52	2.59	**	**	-
	PLACE1003045	16.2	13.12	16.27	11.21	14.83	10.08	8.58	7.66	5.73	**		-
	PLACE1003052	15.26	13.65	16.48	5.57	12.75	6.74	15.39	12.34	13.31	*		-
	PLACE1003083	28.86	32.06	29.46	16.75	26.42	15.92	23.57	20.32	20.14	*		-
	PLACE1003085	11.46	7.5	13.4	5.76	9.36	4.66	6.66	6.58	5.57			
	PLACE1003092	13.44	11.78	13.5	8.16	11.43	7.08	8.07	7.87	6.9	*	**	-
50	PLACE1003097	6.17	6.46	16.32	3.05	5.89	3.08	3.13	5.88	4.77			
	PLACE1003100	36.23	40.39	38.56	32.04	32.87	28.75	26.62	23.05	18.02	*	**	-
	PLACE1003108	11.34	8.75	20.83	7.36	13.72	5.15	3.96	4.75	4.82			
	PLACE1003115	7.34	6.75	7.41	3.32	6.32	2.75	1.65	0.73	1.92	**		-
	PLACE1003120	12.49	11.78	16.03	7.63	9.21	4.76	11.87	11.32	7.77	*		-
	PLACE1003135	7.09	5.05	7.89	3.8	2.81	2.79	2.64	3.42	1.44	*	*	-
55	PLACE1003136	68.81	77.15	81.76	65.96	72.38	60.39	44.74	39.52	47.1	**		-
	PLACE1003141	9.02	4.98	11.12	4.59	4.76	5.17	4.62	5.32	3.84			

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	PLACE1002140	8.54	9.5	15.02	4.83	8.98	6.88	2.91	6.38	4.28	*	-
	PLACE1002150	27.18	33.61	30.05	18.28	25.09	23.98	12.7	14.95	15.29	* **	- -
5	PLACE1002153	173.25	217.95	284.37	422.44	424.65	291.99	262.39	242.27	220.74	*	+
	PLACE1002157	7.42	5.99	8.84	3.16	6.59	2.52	3.59	2.5	5.34	*	-
	PLACE1002163	9.5	10.33	17.61	4.81	9.35	4.71	6.61	8.07	8.46	*	-
	PLACE1002168	406.19	407.62	562.52	187.65	80.86	161.78	377.56	231.11	429.49	**	-
	PLACE1002170	23.07	18.68	22.02	9.58	14.95	10.12	4.39	9.86	5.14	* **	- -
	PLACE1002171	118.3	86.27	82.5	107.58	135.49	96.73	60.2	59.88	56.63	*	-
10	PLACE1002180	12.53	12.02	17.4	11.66	15.77	12.8	11.61	13.11	10.84	*	-
	PLACE1002184	76.94	80.34	67.89	42.12	70.86	43.23	20.87	27.8	29.07	**	-
	PLACE1002200	5.64	6.44	11.56	2.73	4.37	3.29	2.8	2.86	3.48	*	-
	PLACE1002205	5.47	5.6	9.09	2.28	6.05	2.28	3.51	2.6	1.9	*	-
	PLACE1002213	11	9.68	7.88	3.37	8.5	3.33	5.73	2.26	4.33	*	-
	PLACE1002219	30.67	24.88	33.1	11.7	18.03	17.11	5.85	10.58	12.01	* **	- -
15	PLACE1002227	5.38	6.1	6.51	3.19	5.95	3.48	3.48	8.52	2.37	*	-
	PLACE1002253	7.93	4.46	10.74	3.51	5.52	2.84	4.18	3.76	3.24	*	-
	PLACE1002256	10.19	4.49	11.42	3.66	5.44	3.17	1.81	4.48	5.21	*	-
	PLACE1002259	6.68	9.14	16.04	2.32	8.56	3.26	2.76	5.45	13.61	*	-
	PLACE1002285	15.21	20.2	16.69	10.84	18.4	10.19	8.14	7.95	6.12	**	-
20	PLACE1002301	11.84	7.92	13.87	3.5	5.75	4.91	3.98	8.04	4.04	*	-
	PLACE1002310	9.3	8.21	7.36	1.56	5.25	3.56	3.57	2.81	3.34	* **	- -
	PLACE1002311	6.47	4.93	7.42	3.5	5.65	2.75	2.49	2.97	1.32	*	-
	PLACE1002319	23.43	17.43	19.83	7.96	14.77	10.74	10.78	13.23	11.96	*	-
	PLACE1002329	7	4.89	6.25	3.06	3.73	2.65	1.68	2.85	1.64	* **	- -
	PLACE1002333	16.35	14.3	13.78	9.29	11.1	10.53	8.49	6.32	5.51	** **	- -
25	PLACE1002342	14.18	15.16	20.35	5.96	11.78	6.24	4.58	8.15	6.84	* **	- -
	PLACE1002343	9.54	8.56	16.64	3.45	4.23	3.21	2.86	1.85	3.18	*	-
	PLACE1002355	8.47	11.34	12.23	4.32	8.27	5.65	2.98	2.78	4.19	* **	- -
	PLACE1002358	6.25	5.01	7.85	2.33	4.09	1.66	2.63	1.27	2.04	* **	- -
	PLACE1002359	94.59	64.3	78.26	42.47	26.68	29.68	51.17	53.76	59.14	**	-
	PLACE1002374	11.64	7.08	11.73	4.84	7	4.57	3.23	3.62	3.66	*	-
30	PLACE1002376	10.84	6.42	7.57	6.56	6.2	4.93	8.79	6.04	6.62	*	-
	PLACE1002379	14.26	10.82	12.17	6.95	8.14	6.98	5.34	4.44	3.6	** **	- -
	PLACE1002386	7.89	3.52	7.76	1.91	2.26	2.95	2.4	2.03	2.94	*	-
	PLACE1002395	17.68	20.65	22.77	6.45	8.6	7.04	4.3	2.63	2.39	** **	- -
	PLACE1002399	9.59	8.59	10	4	5.08	4.5	2.83	2.8	3.46	** **	- -
	PLACE1002407	14.84	17.54	12.15	4.78	9.29	6.64	5.32	3.95	6.87	* **	- -
35	PLACE1002433	18.89	14.68	17.11	8	12.47	6.81	12.74	12.61	16.41	*	-
	PLACE1002437	6.94	5.04	7.88	4.77	3.72	3.8	1.97	3.5	4.81	*	-
	PLACE1002438	8.47	9.76	8.93	6.11	6.03	2.52	3.69	3.74	4.55	* **	- -
	PLACE1002446	61.22	60.91	53.35	35.08	40.23	34.18	52.04	54.81	48.63	**	-
	PLACE1002447	50.06	38.25	48.75	30.92	38.93	28.56	26.38	23.5	18.47	**	-
40	PLACE1002450	8.68	4.18	8.14	2.56	3.07	3.06	3.45	3.27	1.66	*	-
	PLACE1002462	20.62	21.36	20.69	8.83	13.93	7.63	14.57	11.43	17.67	** *	- -
	PLACE1002465	6.3	10.21	4.45	2.37	6.62	4.25	3.41	1.81	3.99	*	-
	PLACE1002474	8.9	10.86	8.01	5.33	8.28	3.73	2.36	3.29	4.22	**	-
	PLACE1002477	10.23	12.91	13.31	8.45	8.55	11.89	12.5	6.6	14.96	*	-
	PLACE1002493	5.44	5.3	4.56	4.99	2.63	4.56	5.81	1.5	1.86	*	-
45	PLACE1002497	9.23	6.42	6.03	5.54	3.94	3.18	4.77	3.01	2.42	*	-
	PLACE1002499	28.78	23.91	22.93	13.59	10.19	8.79	16.45	7.7	12.71	** *	- -
	PLACE1002500	9.83	6.14	9.08	4.34	6.18	2.49	5.4	1.41	3.42	*	-
	PLACE1002514	57.39	61.86	51.39	31.16	36.96	26.56	53.37	39.47	40.85	**	-
	PLACE1002518	7.97	7.12	4.14	3.92	5.36	3.52	3.58	1.82	2.65	*	-
	PLACE1002529	14.35	10.84	8.84	6.09	12.13	6.95	3.37	3.6	3.7	**	-
	PLACE1002532	6.92	6.97	5.35	4.94	5.47	4.76	4.62	5.36	3.19	*	-
50	PLACE1002536	9.3	12.32	10.38	5.35	4.64	5.41	7.12	5.91	4.59	** *	- -
	PLACE1002537	7.11	8.85	6.37	5.99	5.45	6.3	4.25	3	3.53	**	-
	PLACE1002539	7.92	6.45	7.28	2.97	5.46	7.97	2.07	2.63	4.82	*	-
	PLACE1002547	14.65	14.48	12.7	9.78	11.89	6.33	5.01	4.47	6.38	**	-
	PLACE1002571	27.85	40.71	17.49	16.84	14.71	12.01	13.72	7.48	9.31	*	-
	PLACE1002578	11.7	10.9	10.03	8.42	11.59	6.31	11.22	5.24	9.68	*	-
55	PLACE1002583	12.45	8.98	8.93	4.7	6.58	5.37	8.92	7.31	10.13	*	-
	PLACE1002591	14.95	9.64	30.65	14.99	14.6	1.99	10.65	10.99	7.37	*	-
	PLACE1002598	134.77	135.3	115.7	105.05	127.67	81.8	77.51	75.71	89.77	**	-

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	PLACE1001716	13.89	6.26	12.85	5.68	7.61	3.53	6.17	4.09	5.84				
	PLACE1001720	20.23	23.86	21.86	17.41	19.42	17.41	10.68	9.34	8.72	*	**	-	-
	PLACE1001728	21.97	14.68	15.36	7.87	10.67	10.96	10.39	5.46	5.25	*	*	-	-
5	PLACE1001729	7.28	8.78	9.32	4.24	3.8	3.29	3.96	2.03	2.64	**	**	-	-
	PLACE1001739	18.08	18.75	18.08	11.34	16.38	11.72	10.55	8.26	8.75	*	**	-	-
	PLACE1001740	60.6	61.56	51.85	52.21	47.26	48.39	41.09	28.81	37.51		**	-	-
	PLACE1001745	155.43	121.46	146.04	80.69	92.15	78.37	106.17	102.99	100.25	**	*	-	-
	PLACE1001746	12.07	9.53	11.89	5.08	6.32	6.38	6.4	7.42	7.33	**	**	-	-
10	PLACE1001748	20.09	15.85	13.57	8.98	10.94	8.77	6.58	3.28	8.51	*	*	-	-
	PLACE1001753	8	5.61	7.17	2.42	2.42	2.08	3.13	1.99	2.1	**	**	-	-
	PLACE1001760	16.81	13.89	14.09	5.89	6.99	11.7	8.68	6.91	7.81	*	**	-	-
	PLACE1001767	7.75	5.83	9.04	3.14	5.58	2.6	2.92	2.72	1.2	*	**	-	-
	PLACE1001771	6.86	8.78	6.5	5.1	22.6	7.92	6.68	3.61	1.79			-	-
	PLACE1001775	10.25	8.72	8.89	5.62	4.98	4.94	5.79	4.51	5.33	**	**	-	-
15	PLACE1001777	55.87	33.24	46.38	19.69	26.37	22.71	26.45	22.89	29.83	*		-	-
	PLACE1001781	7.62	4.99	5.26	2.61	3.65	3.27	2.46	2.47	3	*	*	-	-
	PLACE1001783	8.67	3.54	4.17	1.75	1.71	2.1	2.79	1.87	2.94			-	-
	PLACE1001786	10.13	7.03	6.19	5.28	7.53	4.48	4.3	2.57	2.1	*		-	-
	PLACE1001788	9.62	8.2	7.79	2.13	3.02	3.67	3.87	1.93	2.04	**	**	-	-
	PLACE1001795	12.86	13.57	11.91	3.76	6.82	4.77	2.63	4.68	2.65	**	**	-	-
20	PLACE1001799	15.19	12.46	11.73	6.44	10.22	5.84	11.66	9.72	12.14	*		-	-
	PLACE1001810	19.9	17.99	15.3	8.61	16.81	15.9	19.42	15.41	19.52			-	-
	PLACE1001817	9.2	10.03	10.24	5.73	12.6	6.09	4.49	4.6	5.41		**	-	-
	PLACE1001821	8.12	9.65	9.24	5.34	5.81	5.22	5.09	4.7	3.49	**	**	-	-
	PLACE1001836	14.74	10.74	8.7	7.59	8.81	4.93	6.55	5.02	5		*	-	-
	PLACE1001844	8.12	5.11	7.16	1.96	5.82	2.85	3.18	2.67	1.5	*		-	-
25	PLACE1001845	13.24	11.82	8.77	7.59	7.48	3.74	4.84	2.2	2.42	**		-	-
	PLACE1001858	9.06	7.62	7.23	4.16	3.93	3.22	7.66	3.49	3.91	**		-	-
	PLACE1001869	5.88	6.08	4.84	3.37	6.33	2.79	3.14	2.33	2.54		**	-	-
	PLACE1001880	14.71	9.13	7.02	7.27	10.27	7.92	6.48	7.41	7.57			-	-
	PLACE1001887	21.63	13.66	12.13	7	10.58	7.47	6.17	7.42	7.81	*		-	-
	PLACE1001902	25.9	24.36	23.51	11.89	19.02	13.01	23.32	19.87	28.51	*		-	-
30	PLACE1001904	4.8	4.58	4.71	4.2	4.02	2.48	4	1.2	3.18			-	-
	PLACE1001907	11.99	11.09	12.77	4.36	7.42	4.55	5.4	2.81	6.4	**	**	-	-
	PLACE1001910	17.55	13	11.78	10.79	10.86	3.78	8.75	3.34	6.51	*		-	-
	PLACE1001912	16.99	20.76	18.01	10.56	10.89	7.91	12.99	8.39	12.1	**	*	-	-
	PLACE1001918	10.35	11.16	9.4	5.64	8.81	4.76	8.14	4.97	7.28	*	*	-	-
	PLACE1001920	3.62	1.84	1.09	2.06	2.78	2.12	0.89	2.37	2.67			-	-
35	PLACE1001928	33.91	29.31	26.67	16.07	24.12	10.81	21.1	17.01	25.49	*		-	-
	PLACE1001930	13.35	8.41	9.7	5.16	9.65	8.14	5.7	4.65	9.73			-	-
	PLACE1001949	83.24	64.21	68.53	50.5	75.89	66.37	92.93	85.41	92.12	*		-	-
	PLACE1001959	8.04	5.93	10.64	4.56	6.1	3.95	4.33	5.06	5.07			-	-
	PLACE1001969	8.52	5.51	4.9	2.62	4.36	2.04	3.84	2.43	3.2			-	-
40	PLACE1001974	16.8	19.43	17.28	9.77	10.54	5.8	6.88	5.43	6.49	**	**	-	-
	PLACE1001981	11.42	11.82	8.18	7.19	8.78	5.34	7.9	5.15	5.59	*		-	-
	PLACE1001983	9.22	8.71	12.62	5.54	7.95	6.32	7.39	5.63	6.5			-	-
	PLACE1001989	27.47	22.77	28.46	13.3	18.78	12.49	12.35	16.17	10.35	*	**	-	-
	PLACE1002004	8.73	7.48	9.33	6.79	9.77	5.49	6.57	6.13	4.33	*		-	-
	PLACE1002008	10.54	8.97	12.2	6.77	11.46	10.25	3.74	7.33	3.11	*		-	-
45	PLACE1002015	12.37	10.1	13.36	6.64	10.58	8.15	7.89	14.69	9.22			-	-
	PLACE1002044	18.51	15.04	21.28	6.21	11.74	9.02	6.6	10.2	12.21	*	*	-	-
	PLACE1002046	3.16	4.19	5.33	2.46	6.56	3.13	3.41	1.95	2.73			-	-
	PLACE1002052	4.29	2.34	2.47	1.89	4.94	1.55	4.09	3.83	3.62			-	-
	PLACE1002066	4.5	3.49	6.01	2.29	3.79	2.93	3.54	3.65	2.92			-	-
	PLACE1002072	6.4	5.54	6.91	3.66	5.16	4.7	3.95	6.12	2.07	*		-	-
50	PLACE1002073	12.89	11.8	16.36	10.66	11.47	7.97	5.88	5.64	4.74	**		-	-
	PLACE1002080	18.63	22.97	23.1	12.21	18.28	13.9	13.6	15	15.2	*	*	-	-
	PLACE1002081	23.13	27.18	27.3	14.91	18.06	14.86	16.14	20.45	20.82	**	*	-	-
	PLACE1002090	11.49	14.13	12.79	4.54	7.24	4.65	4.29	3.51	5.75	**	**	-	-
	PLACE1002095	5.46	5.54	4.46	5.08	11.47	7.41	4.99	5.27	7.8			-	-
	PLACE1002102	8.22	13.54	14.09	8.11	14.1	7.74	9.04	16.2	14.08			-	-
55	PLACE1002109	17.45	10.93	19.35	12.82	7.15	7.61	9.22	12.81	9.47			-	-
	PLACE1002115	6.48	5.79	9.09	2.86	4.39	3.56	3.32	3.75	3.44	*	*	-	-
	PLACE1002119	3.98	5.58	7.03	2.87	6.47	3.2	3.72	4.12	4.62			-	-

Table 469

	PLACE1001279	20.66	18.57	13.78	9.04	11.1	8.58	6.47	4.72	6	*	**	-	-
	PLACE1001280	13.73	13.22	15.63	11.44	10.47	6.22	11.2	7.22	13.37				
5	PLACE1001294	6.1	5.02	7.03	2.98	5.24	4.37	4.18	2.25	4.36				
	PLACE1001295	9.52	6.34	5.73	4.11	5.38	4.8	4.66	3.72	5.16				
	PLACE1001300	110.16	136.11	96.73	55.88	77.8	69.78	127.86	93.28	123.22	*		-	-
	PLACE1001304	22.83	23.36	19.82	13.09	11.18	12.56	8.64	8.3	14.22	**	**	-	-
	PLACE1001311	22.93	29.08	18.33	13.63	11.98	8.91	18.15	14.72	19.17	*		-	-
	PLACE1001323	5.15	4.73	4.73	2.24	3.77	2.07	2.58	2.19	2.91	*	**	-	-
10	PLACE1001325	6.51	4.36	2.66	3.73	4.01	0.79	3.09	1.19	1.69				
	PLACE1001340	12.71	9.9	7.91	5.93	10.17	3.62	8.49	5.49	7.3				
	PLACE1001344	7.26	5.61	6.3	3.22	4.87	2.35	5.26	3.25	5.47	*		-	-
	PLACE1001351	10.82	9.34	9.06	4.54	4.04	2.83	5.68	5.94	6.17	**	**	-	-
	PLACE1001366	16.46	16.32	12.21	7.96	10.61	8	10.02	6.25	6.08	*		-	-
	PLACE1001377	10.72	11.21	8.12	8.14	11.01	6.18	7.03	5.07	5.95	*		-	-
15	PLACE1001383	8.47	7.74	7.34	7.01	7.31	9.16	4.38	4.51	5.67	**		-	-
	PLACE1001384	5.51	3.13	4.87	2.26	4.64	2.12	1.25	1.77	2.28	*		-	-
	PLACE1001387	3.71	2.66	5.28	2.18	4.13	3.15	1.31	2.35	1.76				
	PLACE1001395	2.97	1.83	2.95	0.9	3.39	0.94	2.26	2.27	0.93				
	PLACE1001399	43.48	36.95	39.47	25.54	39.02	28.91	11.4	22.35	12.37	**		-	-
20	PLACE1001401	8.85	12.13	13.82	4.1	7.84	5.28	5.15	5.58	4.43	*	*	-	-
	PLACE1001407	6.42	8.95	6.88	3.19	5.6	3.04	3.23	2.38	2.93	*	**	-	-
	PLACE1001412	4.65	3.97	3.56	2.52	8.46	2.63	3.1	2.07	4.54				
	PLACE1001414	10.05	7.04	8.23	5.26	7.99	5.53	4.37	7.32	4.41				
	PLACE1001416	18.05	14.76	21.75	8.21	15.69	7.54	12.66	14.94	10.13				
	PLACE1001433	10.35	9.53	10.38	6.47	9.26	5.35	4.33	6.8	5.78	**		-	-
25	PLACE1001440	15.78	8.2	12.06	9.97	10.51	7.97	4.11	5.02	3.17	*		-	-
	PLACE1001456	5.95	5.2	7.6	4.32	4.51	3.68	3.46	6.83	4.09				
	PLACE1001464	7.17	7.13	12.08	4.46	13.26	4.47	2.92	3.59	4.74	*		-	-
	PLACE1001468	5.87	5.89	7.51	4.49	7.19	3.15	4.98	3.89	5.45				
	PLACE1001484	1.93	2.72	3.26	3.17	5.24	2.3	3.27	1.7	4.61				
	PLACE1001500	5.55	5.43	7.47	3.4	4.88	4.91	3.43	5.41	4.22				
30	PLACE1001502	10.13	6.73	10.36	4.95	5.09	3.9	4.24	7.4	4.55	*		-	-
	PLACE1001503	18.94	15.22	23.11	6.62	14.6	9.42	9.19	12.06	11.69		*	-	-
	PLACE1001505	14.33	11.5	15.08	6.94	9.91	8.35	5.05	5.21	4.29	*	**	-	-
	PLACE1001513	9.76	7.84	14.27	4.51	12.01	5.72	4.02	6.18	4.23	*	*	-	-
	PLACE1001516	12.33	10.13	14.29	6.5	8.85	7.91	5.4	6.92	6.78	*	*	-	-
	PLACE1001517	22.78	28.63	26.65	15.6	22.43	12.49	25.55	23.34	19.3				
35	PLACE1001523	4.85	4.2	4.25	1.46	3.64	2.44	3.61	4.37	5.02	*		-	-
	PLACE1001526	15.63	6.8	12.63	5.9	5.99	8.7	6.38	8.18	11.68				
	PLACE1001534	13.26	10.83	12.93	4.83	7.66	4.05	5.45	22.34	4.41	**		-	-
	PLACE1001536	12.35	10.07	11.17	5.95	10.51	7.42	7.47	10.57	7.69				
	PLACE1001545	7.78	9.31	13.04	5.48	7.28	3.94	2.6	3.19	3.02	*		-	-
	PLACE1001551	11.64	13.9	20.49	7.15	9.98	9.21	7.74	8.57	7.76				
40	PLACE1001564	5.33	7.01	11.33	5.64	6.02	6.32	3.9	3.45	6.93				
	PLACE1001570	8.81	7.17	13.9	5.61	7.77	3.89	5.34	9.04	4.82				
	PLACE1001571	30.16	29.44	29.76	18.26	25.49	9.78	12.94	16.78	20.12		**	-	-
	PLACE1001595	49.16	53.02	65.48	26.84	29.86	29.28	20.63	17.19	22.48	**	**	-	-
	PLACE1001602	17.93	20.86	26.59	9.8	15.66	12.03	25.23	43.22	27.97	*		-	-
	PLACE1001603	16	11.85	13.84	6.26	7.5	6.17	10.37	25.06	14.43	**		-	-
45	PLACE1001608	313.79	282.26	299.49	248.62	194.58	212.51	255.55	259.42	224.33	*		-	-
	PLACE1001610	34.83	33.4	50.14	21.2	19.91	21.09	16.25	16.43	15.58	*	*	-	-
	PLACE1001611	16.58	14.57	13.52	8.27	15.57	6.53	5.96	3.97	4.4		**	-	-
	PLACE1001628	15.86	14.27	16.52	6.92	12.73	5.97	9.05	9.4	8.15	*	**	-	-
	PLACE1001632	15.6	14.68	11.95	5.15	11.43	6.76	4.77	3.59	2.95	*	**	-	-
	PLACE1001634	10.26	9.24	12.21	4.71	7.41	4.15	3.93	4.97	4.63	*	**	-	-
50	PLACE1001637	29.75	28.67	36.48	16.79	19.46	16.75	23.41	26.79	22.84	**		-	-
	PLACE1001640	28.95	18.03	24.89	16.93	21.97	12.71	8.91	11.62	9.32	*		-	-
	PLACE1001655	8.24	5.56	6.83	4.09	4.25	2.93	2.13	3.7	1.34	*	*	-	-
	PLACE1001672	8.94	18.15	13.66	4.81	11.29	4.01	5.1	4.86	14.47				
	PLACE1001676	39.58	47.75	47.96	23.16	26.28	21.98	31.5	32.99	28.58	**	**	-	-
	PLACE1001683	7.54	6.2	11.62	2.16	4.9	3.68	2.93	3.43	3.72	*		-	-
55	PLACE1001691	10.4	6.5	11.03	4.05	13.34	5.07	4.28	5.31	2.34	*		-	-
	PLACE1001692	3.74	6.24	4.95	3.58	3.97	3.4	3.88	3.22	2.42				
	PLACE1001705	11.04	7.84	13.66	10.64	5.83	6.51	5.2	5.34	3.62	*		-	-

Table 468

	PLACE1000841	7.87	7.94	8.8	7.6	12.23	10.86	5.33	4.89	4.56	**	-
	PLACE1000843	16.04	20.1	24.44	8.07	13.38	10.99	7.02	9.18	7.59	*	-
5	PLACE1000849	5.43	7.23	9.29	4.99	5.8	4.19	3.26	3.96	2.16	*	-
	PLACE1000856	209.98	229.07	253.03	209.39	256.66	183.75	147.93	119.29	119.6	**	-
	PLACE1000863	16.12	16.52	27.27	10.41	12.7	10.18	13.62	16.56	12.54		-
	PLACE1000876	16.98	21.81	24.49	11.69	18.7	12.24	19.45	15.76	20.21		-
	PLACE1000899	225.37	148.25	253.61	106.96	112.25	68.21	130.08	130.64	120.26	*	-
	PLACE1000907	11.9	9.88	11.62	4.52	6.53	6.56	4.58	7.2	4.58	**	-
10	PLACE1000909	219.81	187.78	224.87	129.92	144.43	97.69	209.39	207.09	231	**	-
	PLACE1000912	4.53	4.31	6.11	1.98	4.16	2.65	2.79	1.25	1.4	*	-
	PLACE1000914	13.65	11.52	16.16	6.76	9.53	7.59	5.22	5.53	2.91	*	-
	PLACE1000918	13.18	11.98	16.82	5.88	9.64	8.52	8.09	14.67	8.16	*	-
	PLACE1000927	6.93	7.22	10.54	4.38	6.32	3.11	4.55	3.47	4.34	*	-
	PLACE1000931	9.5	7.15	10.84	7.43	6.57	4.4	3.12	4.27	3.39	**	-
15	PLACE1000944	12.44	7.29	10.38	5.9	6.11	4.93	5.15	6.79	5.9	*	-
	PLACE1000948	15.03	15	16.76	7.56	11.29	7.62	5.5	9.31	6.83	**	-
	PLACE1000958	5.27	4.99	7.38	2.51	5.7	3.99	1.85	6.23	4.43		-
	PLACE1000972	8.38	5.92	10.49	3.65	7.53	4.25	3.77	5.86	4.19		-
	PLACE1000977	11.29	8.32	15.95	5.04	5.25	7.02	2.5	10.48	3.14		-
	PLACE1000979	13.89	7.86	9.86	3.34	7.43	5.36	4.42	5.93	8.8		-
20	PLACE1000986	30.12	30.53	34.37	25.05	28	23.07	15.94	15.63	18.52	*	-
	PLACE1000987	19.26	23.97	21.74	13.41	21.5	10.91	16.26	13.02	12.47	*	-
	PLACE1001000	4.37	7.15	10.87	3.42	3.04	2.14	2.69	2.93	3.43		-
	PLACE1001007	6.61	2.96	5.44	2.81	6.06	3.04	2.78	3.89	2.07		-
	PLACE1001010	9.46	6.18	10.43	5.02	5.52	3.2	5.18	3.73	3.91	*	-
	PLACE1001015	7.99	5.3	8.03	6.19	7.65	4.6	3.26	3.31	2.3	*	-
25	PLACE1001016	17.51	11.26	16.44	6.61	7.04	-6.42	3.95	3.27	2.38	*	-
	PLACE1001022	14.42	12.9	14.76	4.97	7.37	5.56	7.11	8.21	7.4	**	-
	PLACE1001024	11.18	12.76	12	5.14	10.39	4.2	4.43	3.88	2.99	**	-
	PLACE1001036	105.59	109.4	58.51	52.8	114.22	89.89	113.54	86.02	137.21		-
	PLACE1001038	9.63	8.86	10.53	4.68	5.51	5.33	2.98	5.92	7.98	**	-
	PLACE1001048	7.22	3.94	6.41	5.19	5.21	3.49	3.25	3.01	2.83	*	-
30	PLACE1001054	17.12	18.15	23	21.95	14.11	11.96	6.57	6.82	6.81	**	-
	PLACE1001062	7.69	5.86	7.49	3.03	6.18	4.98	5.6	1.77	1.91	*	-
	PLACE1001063	11.82	13.23	13.09	8.34	8.29	7.45	3.15	3.88	2.78	**	-
	PLACE1001076	17.46	15.27	22.58	9.61	12.22	10.09	9.5	6.93	10.2	*	-
	PLACE1001081	8.04	6.12	10.01	3.34	3.8	3.43	2.11	2.93	3.84	*	-
	PLACE1001088	1077.5	1434.9	897.35	706.08	898.84	874	835.46	1189	1173.1		-
35	PLACE1001092	41.3	35.7	38.59	14.58	20.22	12.39	28.15	30.94	33.71	**	-
	PLACE1001096	27.39	22.29	18.87	5.94	11.11	13.11	8.05	8.36	7.66	*	-
	PLACE1001100	12.04	12.72	14.57	7.69	8.66	8.94	7.95	8.39	9.98	**	-
	PLACE1001104	10.26	8.03	11.29	7.1	4.81	4.72	5.84	4.16	5.12	*	-
	PLACE1001114	19.87	21.11	13.11	5.44	7.05	8.22	2.52	3.44	2.49	*	-
	PLACE1001118	16.04	15.17	13.54	9.49	13.49	10.14	5.69	6.26	3.99	**	-
40	PLACE1001123	11.13	9.28	10.37	3.01	8.71	7.98	2.59	2.22	5.96	**	-
	PLACE1001136	9.63	8.36	10.4	4.16	7.83	10.1	11.4	6.43	4.95		-
	PLACE1001144	22.24	18.89	18.24	5.11	10.59	8.11	16.88	14.03	17.77	**	-
	PLACE1001147	9.73	14.79	11.54	7.5	5.73	6.98	3.33	4.01	4.07	*	-
	PLACE1001148	16.26	12.46	13.41	8.25	12.02	6.54	11.91	10.71	11.91		-
45	PLACE1001159	5.83	2.18	9.14	5.55	4.85	2.75	3.37	5.16	1.28		-
	PLACE1001168	6.64	5.38	5	2.78	4.75	2.95	4.38	2.12	2.36	*	-
	PLACE1001171	17.87	14.63	16.06	9.77	9.8	9.82	9.4	7.26	9.58	**	-
	PLACE1001183	24.74	23.48	20.67	15.23	15.9	11.49	12.71	7.91	11.36	**	-
	PLACE1001185	8.91	7.62	8.02	4.21	7.14	4.11	5.56	3.21	2.94	*	-
	PLACE1001201	16.85	15.12	18.88	10.04	10.01	9	9.33	10.2	8.94	**	-
50	PLACE1001229	15.21	12.61	13.94	9.57	11.26	12.86	8.81	8.47	8.63	**	-
	PLACE1001231	13.31	19.61	12.2	8.58	26.11	14.91	13.39	13.43	8.59		-
	PLACE1001238	6.16	6.8	5.78	3.46	6.45	5.07	3.57	1.84	3.63	**	-
	PLACE1001241	7.81	12.55	8.77	4.56	8.8	2.29	6.75	1.69	5.45		-
	PLACE1001242	15.15	9.43	9.57	6.93	8.62	5.05	5.42	2.86	4.67	*	-
	PLACE1001247	18.19	25.95	17.43	7.95	13.13	8.09	9.19	7.53	8.42	*	-
55	PLACE1001250	27.19	16.6	15.62	14.1	21.99	16.23	13.87	8.35	7.98		-
	PLACE1001257	5.07	4.07	3.3	6.21	3.6	2.05	3.64	3.01	2.62		-
	PLACE1001272	11.61	5.34	6.33	4.73	5.73	6.35	5.18	6.02	6.39		-

Table 467

	PLACE1000374	4.36	4.25	6.34	4.3	3.67	2.84	1.94	3.1	1.77	*	*	-
	PLACE1000380	16.1	6.91	17.82	8.8	10.53	4.49	5.18	5.02	6.92	*	*	-
5	PLACE1000383	6.36	4	6.16	2.37	2.92	2.52	2.89	0.95	0.87	*	*	-
	PLACE1000397	11.59	7.26	11.9	2.79	5.08	5.01	6.09	6.59	6.48	*	*	-
	PLACE1000401	9.04	6.59	7.42	3.23	3.26	3.54	3.28	2.6	2.94	**	**	-
	PLACE1000406	13.08	8.33	14.95	2.59	5.95	3.66	6.86	10.26	6.27	*	*	-
	PLACE1000412	8.51	7.48	7.35	3.42	4.96	3.44	3.84	4.02	2.52	**	**	-
	PLACE1000420	18.93	23.33	27.28	7.24	14.78	9.32	4.07	6.78	6.75	*	**	-
10	PLACE1000421	17.83	12.09	14.21	5.67	5.38	4.01	4.24	5.77	6.73	**	**	-
	PLACE1000423	9.91	10.66	9.25	5.17	5.46	3.16	3.23	3.33	2.82	**	**	-
	PLACE1000424	6.41	4.46	8.86	3.52	3.38	1.84	2.63	2.91	2.34	*	*	-
	PLACE1000430	9.9	7.48	10.47	5.49	5	3.45	3.57	3.09	3.55	*	**	-
	PLACE1000433	8.1	6.1	7.11	1.95	4.44	2.14	2.34	1.88	2.11	*	**	-
	PLACE1000435	28.95	26.53	24.32	14.09	14.79	9.09	16.84	11.65	12.23	**	**	-
15	PLACE1000437	13.7	9.76	13.69	6.32	11.96	9.08	8.69	6.76	4.42	*	*	-
	PLACE1000442	4.21	4.33	5.54	1.97	3.93	1.28	2.62	4.3	2.3			-
	PLACE1000444	8.39	8.02	5.47	3.23	8.73	7.32	3.64	5.13	5.23			-
	PLACE1000453	8.24	8.08	7.4	2.64	6.34	3.96	5.87	6.05	5.1	*	**	-
	PLACE1000456	9.14	8.93	7.88	4.6	6.83	3.55	4.93	5.65	3.15	*	**	-
	PLACE1000455	29.59	27.02	24.71	10.66	16.38	9.53	7.25	7.03	4.61	**	**	-
20	PLACE1000481	10.19	8.45	9.18	6.24	5.2	4.46	5.55	2.41	2.95	**	**	-
	PLACE1000492	14.54	13.99	15.18	6.98	17.38	5.81	5.24	4	6.05	**	**	-
	PLACE1000508	8.74	6.29	6.02	2.46	6.75	1.48	4.68	3.01	3.04	*	*	-
	PLACE1000512	4.85	3.74	3.65	2.37	3	3.01	2.27	2.44	1.75	*	*	-
	PLACE1000540	12.96	8.47	7.79	4.5	9.85	7.03	5.75	9.5	3.54			-
25	PLACE1000541	10.16	15.98	11.44	6.25	17.26	10.37	7.65	6.38	9.85			-
	PLACE1000546	13.1	11.96	11.32	7.18	10.25	10.48	6.08	5.2	6.25	**	*	-
	PLACE1000547	8.49	5.52	7.86	5.49	10.6	14.98	5.43	2.62	3.64	*	*	-
	PLACE1000560	22.37	15.31	18.14	8.36	10.85	7.86	6.26	3.35	3.9	*	**	-
	PLACE1000582	8.22	7.18	9	4.62	4.33	3.02	4.18	2.13	2.7	**	**	-
	PLACE1000584	17.41	14.24	11.35	10.42	10.44	10.34	8.72	6.17	6.09	*	*	-
30	PLACE1000583	11.44	8.04	7.49	4.72	4.9	3.91	5.24	5.95	5.26	*	*	-
	PLACE1000587	7.15	6.59	3.7	3.14	3.88	4.15	3.78	3.64	4.07			-
	PLACE1000588	18.91	11.92	9.8	5.65	7.81	8.36	8.6	5.83	6.87			-
	PLACE1000596	11.7	13.56	11.65	8.47	13.84	9.08	10.58	5.53	9.44			-
	PLACE1000599	9.76	11.08	9.94	6.95	10.5	6.09	4.81	2.43	3.16	**	*	-
	PLACE1000605	16.65	12.34	15.55	9.49	9.68	6.14	6.32	3.07	5.49	*	**	-
35	PLACE1000610	13.13	11.57	11.01	8.74	7.05	4.41	9.69	6.2	4.65	*	*	-
	PLACE1000611	44.34	42.43	32.52	26.13	31.67	19.73	63.56	39.23	50.7			-
	PLACE1000626	7.16	6.36	6.63	3.54	3.14	2.93	5.25	5.57	5.32	**	**	-
	PLACE1000632	7.75	6.07	5.61	4.09	5.95	4.33	4.51	3.72	5.54			-
	PLACE1000636	9.18	4.57	3.73	3.4	5.99	3.07	2.89	4.22	5.33			-
	PLACE1000653	13.52	15.36	9.28	7.56	11.63	7.5	6.22	4.09	7.58	*	*	-
40	PLACE1000656	16.45	24.27	20.78	6.6	13.98	8.95	13.7	10.63	9.13	*	*	-
	PLACE1000663	17.59	14.23	11.72	6.71	8	7.43	15.75	11.07	16.5	*	*	-
	PLACE1000706	22.58	27.08	16.29	14.32	16.15	9.77	10.6	7.13	11.87	*	*	-
	PLACE1000712	15.91	18.32	12.16	10.41	18.03	9.89	15.21	13.33	14.12			-
	PLACE1000716	9.51	6.38	10.32	4.38	8.28	5.91	5.52	6.18	6.28			-
	PLACE1000740	62.65	63.87	75.82	66.13	88.56	67.41	54.02	47.28	39.42	*	*	-
45	PLACE1000748	21.69	20.13	21.77	15.61	19.67	14.26	16.35	17.23	19.62	*	*	-
	PLACE1000749	8.09	6.77	9.49	5.21	7.14	4.97	2.01	3.41	3.49			-
	PLACE1000751	5.85	6.89	7.76	3.42	4.79	3.39	2.03	3.31	2.82	*	**	-
	PLACE1000755	17.27	23.58	23.84	13.28	21.64	9.26	15.55	10.98	9.19	*	*	-
	PLACE1000769	3.3	5.09	3.02	3.66	5.56	3.4	3.76	2.68	2.93			-
	PLACE1000772	43.25	43.09	44.83	22.77	18.03	27.54	20.87	24.08	21.65	**	**	-
50	PLACE1000785	31.35	17.11	31.03	9.35	9.07	8.25	11.86	8.82	7.46	*	*	-
	PLACE1000786	5.99	3.18	7.12	2.4	4.01	1.88	2.45	4.94	6.92			-
	PLACE1000793	62.95	70.84	86.12	74.56	80.1	68.58	59.03	72.61	48.69			-
	PLACE1000795	170.82	165.32	165.81	123.39	175.42	137.1	104.41	82.02	95.27	**	*	-
	PLACE1000798	34.34	30.96	28.31	19.04	23.22	14.81	22.05	22.97	28.25	*	*	-
	PLACE1000812	16.41	17.23	19.15	11.98	12.41	10.24	15.41	13.57	12.8	**	*	-
55	PLACE1000823	25.51	22.51	19.45	15.2	17.94	10.61	13.45	15.88	12.99	*	*	-
	PLACE1000825	9.78	4.9	7.94	5.02	3.89	3.86	4.94	3.55	4.1			-
	PLACE1000838	37.15	32.03	40.12	17.98	18.59	17.35	19.92	15.21	15.97	**	**	-

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	OVARC1002091	18.78	19.82	15.43	10.02	4.76	6.56	9.66	6.62	10.94	**	**	-	-
	OVARC1002092	11.35	7.73	4.3	3.87	5.5	2.76	3.85	4.56	3.79			-	-
5	OVARC1002093	11.71	8.5	20.66	4.96	14.44	7.82	5.85	5.4	7.08			-	-
	OVARC1002094	6.29	6.49	6.65	2.69	4.04	2.24	4.34	3.07	3.08	**	**	-	-
	OVARC1002107	13.48	8.34	12.83	6.06	7.84	6.33	9.41	7.18	12.5	*	*	-	-
	OVARC1002112	26.38	33.01	27.23	12.79	21.35	12.35	12.54	9.82	14.29	*	**	-	-
	OVARC1002126	65.16	69.33	51.38	44.56	51.62	38.84	65.71	52.52	60.89	*	*	-	-
	OVARC1002127	6.15	6.37	5.32	1.95	4.41	3.09	5.22	3.46	3.37	*	*	-	-
10	OVARC1002138	5.22	5.32	3.68	5.16	3.08	2.95	2.56	2.26	2.94	*	*	-	-
	OVARC1002143	7.3	4.6	3.7	3.29	5.14	1.8	3.79	3.64	5.8			-	-
	OVARC1002156	14.33	13.86	13.02	10.03	12.28	6.78	8.59	6.29	5.67	**		-	-
	OVARC1002158	9.15	5.03	4.65	3.18	3.11	2.41	3.88	2.86	4.37			-	-
	OVARC1002165	67.03	61.86	54.25	23.78	36.97	31.97	46.19	44.24	42.15	**	*	-	-
	OVARC1002176	10.61	13.89	9.02	5.22	6.29	4.46	9.03	7.11	8.09	*	*	-	-
15	OVARC1002178	11.02	11.22	6.01	3.16	5.47	2.82	6.17	3.48	3.57	*		-	-
	OVARC1002182	6.32	5.52	5.4	2.92	5.92	3.91	6.46	4.29	5.82			-	-
	OVARC1002185	13.32	14.59	15.01	5.81	10.11	6.86	7.78	7.13	8.82	**	**	-	-
	PLACE1000004	9.74	9.21	11.61	4.85	7.73	5.01	4.83	8.27	4.93	*	*	-	-
	PLACE1000005	27.12	18.01	21.46	13.29	16.54	9.78	14.01	13.75	13.04	*	*	-	-
	PLACE1000006	91.12	57.04	55.23	34.88	52.28	43.96	6.98	11.96	13.8	**	*	-	-
20	PLACE1000007	7.26	8.75	8.28	8.37	10.72	8.06	3.04	5.48	2.7	*		-	-
	PLACE1000014	2.23	2.04	4.15	1.42	4	3.02	1.88	2.43	1.7			-	-
	PLACE1000031	25.49	34.88	32.13	13.67	27.22	11.3	27.49	25.37	24.47			-	-
	PLACE1000033	18.11	15.97	17.95	12.01	14.01	16.72	14.29	10.78	15.77			-	-
	PLACE1000040	8.82	8.21	10.8	5.78	10.45	6.53	3.91	7.12	4.69	*	*	-	-
	PLACE1000048	20.38	17.3	18.51	15.36	23.63	12.54	8.78	9.57	6.66	**	*	-	-
25	PLACE1000050	5.4	7.04	6.5	2.42	4.86	4.46	3.72	6.27	1.7			-	-
	PLACE1000061	9.33	8.61	17.26	5.67	11.39	8.55	3.23	9.76	4.54			-	-
	PLACE1000066	6.66	5.77	17.49	7.42	6.83	5.75	5.92	4.2	3.99			-	-
	PLACE1000075	12.56	13.3	14.47	10.46	23.06	11.81	12.58	8.36	7.76			-	-
	PLACE1000078	16.43	18.48	16.12	10.89	9.95	8.28	13.59	13.8	12.7	**	*	-	-
	PLACE1000081	10.24	9.79	10.46	3.99	5.51	5	3.34	5.78	6.89	**	*	-	-
30	PLACE1000086	24.93	18.79	23.44	11.66	12.45	12.8	14.85	17.83	16.54	**	*	-	-
	PLACE1000094	9.77	8.35	20.14	5.83	7.13	8.45	9.11	6.86	4.47			-	-
	PLACE1000101	13.24	13.69	14.14	7.47	11.68	8.08	3.54	8.15	5.57	*	**	-	-
	PLACE1000121	13.12	18.46	20.42	15.19	20.27	14.52	7.88	10.58	8.77	*	*	-	-
	PLACE1000133	9.91	9.72	12.83	7.9	8.22	8.5	5.09	6.11	8.25	*	*	-	-
35	PLACE1000142	9.88	8.89	10.42	4.19	7.46	3.72	7.05	4.74	5.82	*	**	-	-
	PLACE1000146	26.84	30.96	24.74	12.94	22.4	10.59	30.36	28.2	29.92	*	*	-	-
	PLACE1000163	7.41	5.19	10.36	3.1	5	4.08	3.16	3.43	3.29		*	-	-
	PLACE1000172	27.22	20.45	29.16	10.4	14.2	9.22	8.64	12.45	11.02	**	**	-	-
	PLACE1000181	113.9	71.51	73.22	49.28	64.78	42.83	38.76	53.7	55.02			-	-
	PLACE1000184	17.72	15.27	19.8	11.79	14.01	13.85	9.11	9.87	10.76	*	**	-	-
40	PLACE1000185	16.19	14.48	17.72	7.66	7.1	6.34	4.1	6.59	4.27	**	**	-	-
	PLACE1000198	9.65	11.81	15.71	11.09	11.55	8.4	6.38	19.65	7.03			-	-
	PLACE1000213	6.26	6.05	11.46	4.09	6.36	3.94	3.17	3.29	3.78			-	-
	PLACE1000214	9.15	11.17	11.68	8.73	8.08	4.64	3.79	6.02	5.34		**	-	-
	PLACE1000220	25.57	22.54	23.7	11.03	13.14	7.79	10.18	10.81	12.09	**	**	-	-
	PLACE1000231	14.42	18.43	23.26	5.64	9.79	6.68	7.36	7.88	9.82	*	*	-	-
45	PLACE1000236	7.58	6.02	8.68	1.98	5.24	5.31	3.56	10.92	5.78			-	-
	PLACE1000245	26.52	25.91	22.56	9.19	16.74	15.36	8.75	9.5	10.2	*	**	-	-
	PLACE1000246	45.69	36.58	50.55	14.83	17.16	15.14	8.9	11.47	10.62	**	**	-	-
	PLACE1000258	9.85	11.52	11.33	3.16	7.83	3.08	2.46	5.51	4.37	*	**	-	-
	PLACE1000288	14.3	14.29	20.76	4.5	8.13	8.1	4.73	7.76	5.77	*	*	-	-
	PLACE1000292	22.65	21.9	21.53	9.54	18.93	11.23	12.98	16.24	12.38	*	**	-	-
50	PLACE1000302	24.34	18.15	25.79	8.82	16.47	8.9	10.89	16.25	19.37	*	*	-	-
	PLACE1000304	9.25	6.15	7.74	5.54	8.18	3.68	3.39	11.18	2.7			-	-
	PLACE1000308	15.07	8.66	14.03	6.88	10.38	7.58	8.62	8.08	9.22			-	-
	PLACE1000309	50.33	32.12	45.32	24.2	31.82	23.11	35.7	38.12	29.88			-	-
	PLACE1000312	7.8	7.45	13.99	4.65	6.15	3.95	7.1	6.14	4.22			-	-
	PLACE1000330	37.2	43.13	26.2	23.56	30.84	17.96	20.8	28.35	14.01			-	-
55	PLACE1000332	19.72	22.93	22.77	9.1	18.25	13.71	15.08	10.86	13.37	*	**	-	-
	PLACE1000347	18.65	20.7	25.21	9.08	28.96	11.34	16.06	14.13	12.46	*	*	-	-
	PLACE1000351	4.55	3.12	7.2	2.8	3.17	2.14	6.68	4.21	4.87			-	-

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	OVARC1001702	9.4	5.63	9.29	4.18	2.13	4.1	3.32	3.32	2.46	*	*	-	-
	OVARC1001703	7.15	7.4	12.31	4.47	5.7	4.5	5.85	8.24	3.65	*	*	-	-
5	OVARC1001710	18.66	12.34	27.68	8.37	10.17	11.05	8.89	17.12	13.23	*	*	-	-
	OVARC1001711	18.7	15.2	21.81	10.78	18.26	9.63	7.19	14.6	8.6	*	*	-	-
	OVARC1001713	106.12	95.89	118.84	111.33	127.16	96.1	59.14	62.39	67.07	**	**	-	-
	OVARC1001725	7.41	8.41	15.02	6.01	22.46	3.17	4.32	4.47	6.9	*	*	-	-
	OVARC1001726	9.14	11.76	13.16	5.3	10.91	4.63	7.72	5.13	7.08	*	*	-	-
	OVARC1001727	6.13	4.36	5.44	2.42	3.92	3.64	2.37	4.79	3.09	*	*	-	-
10	OVARC1001731	326.35	314.92	333.21	240.06	116.88	168.02	275.41	235.36	284.79	*	*	-	-
	OVARC1001735	10.45	7.4	10.3	3.26	6.6	5.16	2.85	50.48	3.58	*	*	-	-
	OVARC1001741	51.89	35.06	44.68	30.3	40.05	30.18	29.82	27.49	31.74	*	*	-	-
	OVARC1001745	48.03	48.42	50.14	20.26	24.89	18.25	35.38	41.14	42.19	**	*	-	-
	OVARC1001759	8.58	9.79	14.25	4.9	9.68	2.54	3.21	5.87	3.66	*	*	-	-
	OVARC1001762	15.44	18.76	16.88	4.99	8.2	5.87	5.45	4.57	3.61	**	**	-	-
15	OVARC1001766	83.39	85.91	87.93	59.55	79.17	53.32	53.78	49.12	57.96	**	**	-	-
	OVARC1001767	7.04	5.21	8.42	2.96	5.88	2.78	2.97	1.63	2.74	*	*	-	-
	OVARC1001768	14.52	13.49	18.22	6.31	12.6	7.55	3.97	6.24	6.88	**	**	-	-
	OVARC1001770	26.33	28.28	29.2	13.86	20.54	11.13	6.52	15.1	8.61	*	*	-	-
	OVARC1001776	7.09	4.63	7.55	4.18	5.49	2.76	2.6	3.29	2.71	*	*	-	-
	OVARC1001791	10.39	7.82	10.63	6.37	4.38	3.31	3.58	4.2	5.4	*	**	-	-
20	OVARC1001795	7.98	5.23	12.47	1.59	4.12	3.1	3.31	4.76	7.14	*	*	-	-
	OVARC1001798	40.84	44.26	46	20.27	29.31	16.26	30.23	32.18	29.44	**	**	-	-
	OVARC1001802	21.47	24.32	21.73	9.56	17	10.13	13.8	16.34	14.18	*	**	-	-
	OVARC1001805	21.41	25.19	22.93	13.01	23.64	11.53	6.55	5.13	2.96	**	**	-	-
	OVARC1001807	9.03	4.95	8.69	4.35	5.41	3.66	4.23	6.47	5.73	*	*	-	-
25	OVARC1001809	149.46	129.89	172.57	133.13	197.68	130.62	61.18	73.6	73.85	**	**	-	-
	OVARC1001812	14.81	11.9	18.98	7.06	7.79	6.77	7.18	8.19	7.69	*	*	-	-
	OVARC1001813	16.16	13.4	12.96	5.79	9.88	6.64	9.17	11.83	7.83	*	*	-	-
	OVARC1001820	13.24	13.12	21.66	6.46	7.84	7.81	9.43	11.92	8.26	*	*	-	-
	OVARC1001828	7.36	8.18	15.13	2.53	4.46	3.16	2.97	2.27	1.93	*	*	-	-
	OVARC1001833	7.55	8	10.4	3.73	5.88	2.72	2.14	3.67	2.55	*	**	-	-
30	OVARC1001839	9.51	8.23	9.47	3.48	6.12	2.44	2.33	1.23	3.33	*	**	-	-
	OVARC1001846	10.75	6.47	12.96	4.44	5.49	4.43	2.19	5.18	4.87	*	*	-	-
	OVARC1001849	21.23	22.64	18.18	12.84	14.77	8.26	9.88	17.09	10.17	*	*	-	-
	OVARC1001861	14.15	12.35	14.96	7.86	9.2	6.06	6.33	6.39	8.07	**	**	-	-
	OVARC1001873	10.37	9.13	10.41	3.08	3.81	3.28	3.75	2.72	3.55	**	**	-	-
	OVARC1001879	12.35	13.47	14.5	4.79	6.62	4.85	9.81	8.26	4.82	**	*	-	-
35	OVARC1001880	17.95	16.04	17.13	6.7	12.3	6.25	9.7	7.19	8.89	*	**	-	-
	OVARC1001883	10.68	7.64	9.62	5.66	5.31	4.96	4.5	5.37	3.66	*	**	-	-
	OVARC1001900	21.57	23.04	21.79	9.35	13.16	8.64	4.93	5.9	5.98	**	**	-	-
	OVARC1001901	6.1	3.23	6.27	3.76	3.35	2.44	2	2.36	2.55	*	*	-	-
	OVARC1001911	5.85	6.08	5.24	3.87	3.1	3.23	3.29	4.37	2.41	**	*	-	-
	OVARC1001916	15.55	14.04	13.05	10.21	8.26	9.44	6.7	7.73	11.27	**	*	-	-
40	OVARC1001928	4.28	4.17	5.98	2.82	2.37	1.14	3.65	1.6	1.43	*	*	-	-
	OVARC1001937	25.6	15.71	13.44	9.48	11.18	11.27	11.49	12.97	7.51	*	*	-	-
	OVARC1001940	8.76	8.78	8.37	2.83	3.28	2.84	2.16	2.96	2.02	**	**	-	-
	OVARC1001942	20.24	15.16	17.14	7.55	9.31	5.48	5.3	6.71	7.34	**	**	-	-
	OVARC1001943	29.68	32.41	20.55	12.97	22.62	15.27	4.34	4.72	5.72	**	**	-	-
	OVARC1001949	14.19	11.75	10.33	3.19	5.85	4.46	2.89	6.47	5.47	**	**	-	-
45	OVARC1001950	13.94	15.9	10.94	6.19	9.29	6.75	8.04	7.58	9.81	*	*	-	-
	OVARC1001952	122.9	130.05	103.73	90.55	135.58	102.89	79.11	70.36	72.15	**	**	-	-
	OVARC1001954	7.45	6.81	6.15	3.11	3.63	2.91	2.32	2.33	2.18	**	**	-	-
	OVARC1001963	12.44	11.79	17.61	6.04	6.32	5.28	6.38	9.47	7.04	*	*	-	-
	OVARC1001983	25.71	30.08	25.81	12.15	15.68	10.47	16.02	11.31	11.87	**	**	-	-
	OVARC1001987	13.81	15.75	12.25	6.15	7.8	4.37	5.37	6.61	4.93	**	**	-	-
50	OVARC1001989	18.41	16.89	13.96	7.74	13.38	8.71	14.94	11.39	11.29	*	*	-	-
	OVARC1001991	12.31	10.21	10.74	6.47	8.3	7.41	6.69	6.19	5.31	*	**	-	-
	OVARC1002005	33.83	38.39	35.62	17.89	27.06	20.01	28.16	23.73	27.11	**	**	-	-
	OVARC1002044	25.13	24.53	23.58	14.08	17.38	11.27	23.29	17.04	24.49	**	**	-	-
	OVARC1002046	79.06	107.27	86.51	49.53	77.97	65.8	68.65	65.46	67.43	*	*	-	-
	OVARC1002050	9.22	6.62	10.19	4.05	4.76	4.15	4.13	2.81	4.5	*	*	-	-
55	OVARC1002058	18.18	16.91	17.55	13.19	18.87	6.99	24.17	22.71	28.4	*	*	-	-
	OVARC1002068	67.02	72.72	47.65	37.79	39.59	35.74	62.17	50.77	51.33	*	*	-	-
	OVARC1002082	34.4	33.12	24.78	17.13	27.63	15.59	48.32	29.05	33.16	*	*	-	-

Table 464

	OVARC1001268	91.7	87.42	103.18	50.09	61.38	43.24	78.98	90.56	56.28	**	-
	OVARC1001270	7.04	6.65	7.37	4.89	4.4	3.37	2.36	3.78	2.58	**	*
5	OVARC1001271	19.59	14.31	19.8	9.15	12.99	7.59	10.4	9.45	13.28	*	*
	OVARC1001282	9.08	5.35	8.29	3.47	3.89	2.86	2.79	2.02	1.62	*	*
	OVARC1001296	8.37	7.88	7.76	3.73	3.83	3.51	3.2	2.52	1.42	**	**
	OVARC1001306	8.13	8.36	10.06	2.51	4.13	1.61	1.91	3.24	2.11	**	**
	OVARC1001314	6.35	7.88	10.42	5.15	6.88	5.6	5.8	2.73	3.16	*	*
	OVARC1001316	9.21	9.19	7.4	3.28	5.15	2.41	2.09	2	2.6	**	**
10	OVARC1001329	66.76	60.97	62.75	30.71	36.83	28.85	49.5	54.71	56.06	**	*
	OVARC1001330	6.94	7.09	6.08	3.83	3.92	2.72	1.52	3.71	2.62	**	**
	OVARC1001336	20.53	13.52	17.21	8.37	10.93	7.02	7.67	13.22	12.66	*	*
	OVARC1001338	9.67	3.35	7.39	3.31	2.87	2.49	3.22	2.79	2.03	*	*
	OVARC1001339	93.84	87.95	84.06	45.91	66.98	50.22	43.37	46.73	40	**	**
	OVARC1001340	6.83	5.17	8.34	3.19	3.73	1.7	2.64	3.21	3.16	*	*
15	OVARC1001341	30.38	40.21	33.21	15.81	17.56	16.77	23.98	16.89	14.16	**	*
	OVARC1001342	319.96	384.64	298.02	528.73	514.44	449.66	353.76	251.2	152.62	*	+
	OVARC1001344	24.05	19.69	20.93	8.91	10.1	7.96	16.66	12.81	15.35	**	*
	OVARC1001357	6.41	18.75	6.59	4.08	6.56	3.73	3.75	4.14	3.16	*	*
	OVARC1001359	59.32	54.06	31.23	23.17	43.75	45.21	51.59	26.81	37.78	*	*
20	OVARC1001360	5.67	4	3.9	1.82	2.67	1.87	1.59	2.07	1.71	*	**
	OVARC1001369	7.6	9.8	9.35	5.24	6.16	2.48	5.5	3.41	1.68	*	*
	OVARC1001372	9.97	8.14	5.06	1.49	5.79	2.2	2.9	3.07	0.96	*	*
	OVARC1001376	27	25.34	21.25	12.73	15.31	10.9	17.23	14.16	15.91	**	*
	OVARC1001381	39.84	52.86	35.2	18.78	31.21	17.84	39.83	32.09	35.47	*	*
	OVARC1001391	4.35	4.74	4.59	3.41	5.61	3.07	2.93	3.15	1.97	*	**
25	OVARC1001392	11.01	13.87	11.26	7.11	8.72	6.36	4.32	4.97	3.08	*	**
	OVARC1001399	7.11	9.27	5.73	3.62	4.78	3.62	4.04	3.29	4.7	*	*
	OVARC1001417	8.65	8.11	7.05	3.69	5.18	3.15	2.59	2.11	3.11	**	**
	OVARC1001419	9.55	7.34	7.59	3.69	5.75	3.4	4.52	2.92	1.58	*	**
	OVARC1001425	13.5	5.66	6.59	3.04	5.54	5.43	3.66	1.31	1.7	*	*
	OVARC1001436	14.91	13.09	11.35	5.12	5.94	3.89	4.56	3.69	3.48	**	**
30	OVARC1001442	5.57	6.26	6.22	3.11	6.56	2.54	5.62	2.71	2.72	*	*
	OVARC1001451	30.68	20.35	20.93	10.57	21.04	10.84	19.07	21.71	23.53	*	*
	OVARC1001452	16.95	14.67	11.44	7.52	6.15	5.24	7.7	11.36	6.64	**	*
	OVARC1001453	7.86	6.06	6.61	4.22	6.52	4.39	4.72	4.86	4.79	*	*
	OVARC1001476	20.63	24.57	30.44	15.92	22.32	7.34	11.05	12.16	16.07	*	*
	OVARC1001480	7.35	4.8	7.98	3.71	4.84	5.16	2.87	2.08	3.16	*	*
35	OVARC1001489	11.25	5.88	6.53	8.16	19.5	4.38	4.69	2.8	2.81	*	*
	OVARC1001493	15.28	11.36	13.67	4.88	4.94	2.76	9.11	6.21	8.31	**	*
	OVARC1001496	32.05	27.24	23.8	12.75	20.64	11	9.38	5.04	10.85	*	**
	OVARC1001499	5.67	6.05	4.7	2.72	3.78	1.85	2.85	4.6	2.57	*	*
	OVARC1001506	31.76	25.87	22.22	8.57	18.43	9.44	18.79	15.06	20.59	*	*
	OVARC1001509	17.78	13.64	12.09	7.06	8.79	5.76	15.13	14.75	21.14	*	*
40	OVARC1001510	8.59	5.24	6.08	3.07	5.95	3.22	7.27	7.67	4.5	*	*
	OVARC1001516	10.81	12.88	13.12	5.8	9.08	6.34	8.79	7.64	9.61	*	*
	OVARC1001525	7.75	5.51	6.34	3.13	4.32	3.66	4.28	3.77	3.8	*	*
	OVARC1001542	30.12	29.23	28.39	14.89	25.03	14.49	18.26	13.2	26.28	*	*
	OVARC1001544	27.99	28.47	20.46	12.55	18.48	10.77	36.56	24.05	33.39	*	*
	OVARC1001546	6.07	8.98	10.22	3.38	5.35	8.68	3.38	3.45	4.28	*	*
45	OVARC1001547	4.36	6.21	6.19	3.96	3.95	1.63	2.77	2.7	2.24	**	*
	OVARC1001555	137.04	110.6	136.25	120.36	174.1	119.31	90.66	69.83	70.87	*	*
	OVARC1001560	10.54	9.21	9.05	6.2	10.38	4.19	4.33	6.13	2.34	*	*
	OVARC1001569	28.34	25.69	28.31	17.71	19.82	21.58	15.15	17.9	15.82	**	**
	OVARC1001570	9.57	12.59	13.54	7.21	9.82	6.14	6.01	6.08	8.45	*	*
	OVARC1001577	10.2	13.78	13.09	13.59	16.75	9.11	10.56	8.87	5.25	*	*
50	OVARC1001578	0	0.27	1.12	0.09	2	0.48	1.37	0.77	0	*	*
	OVARC1001596	18.44	38.66	31.49	14.48	23.49	16.88	13.34	13.82	12.27	*	*
	OVARC1001600	12.84	12.05	12.79	8.39	9.83	7.52	7.16	10.38	7.31	**	*
	OVARC1001607	17.69	15.72	17.94	10.9	18.71	12.05	10.18	16.33	11.6	*	*
	OVARC1001610	7.42	7.24	11.17	5.09	7.94	4.34	2.78	4.22	3.55	*	*
	OVARC1001611	3.52	5.78	9.33	2.82	5.44	3.47	2.68	4.9	2.76	*	*
55	OVARC1001615	7.22	4.15	9.71	3.66	4.51	3.88	4.88	4.91	5.78	*	*
	OVARC1001636	5.13	4.74	6.26	4.34	6.6	4.17	6.58	6.24	9.12	*	*
	OVARC1001668	40.93	55.07	40.85	20.99	33.29	18.89	40.58	43.41	37.28	*	*

Table 463

	OVARC1000996	12.75	15.42	13.22	7.58	7.82	4.86	10.5	9.45	10.8	**	*	-	-
	OVARC1000999	77.92	64.11	52.33	34.61	41.57	26.78	51.13	38.47	53.84	*	-	-	-
5	OVARC1001000	34.25	25.88	23.55	13.32	17.72	11.08	24.85	23.48	28.11	*	-	-	-
	OVARC1001004	4.45	4.47	4.74	3.56	4.7	2.11	3.94	2.09	3.24				
	OVARC1001010	7.8	4.93	6.92	2.94	3.81	2.97	3.34	2.1	2.69	*	*	-	-
	OVARC1001011	10.56	5.96	7.28	2.55	4.47	3.01	5.82	4.52	6.66	*	-	-	-
	OVARC1001030	135.15	261.75	227.94	119.93	118.81	67.28	114.11	94.32	117.34				
	OVARC1001032	10.53	7.53	10.42	4.6	4.01	3.78	8.13	6.94	7.33	**	-	-	-
10	OVARC1001034	10.83	11.24	8.27	5.28	4.2	4.55	7.53	5.89	7.35	**	*	-	-
	OVARC1001038	36.07	37.32	30.24	14.88	24.71	11.8	14.41	10.88	16.24	*	**	-	-
	OVARC1001040	40.95	33.02	36.27	20.4	24.74	15.2	38.11	31.08	34.73	**	-	-	-
	OVARC1001041	28.88	24.51	24.41	14.12	19.17	10.91	20.98	17.34	23.83	*	-	-	-
	OVARC1001044	8.66	8.02	6.34	3.74	2.76	3.79	6.52	3.05	4.17	**	-	-	-
	OVARC1001049	23.67	22.64	22.97	8.16	12.73	9.28	19.46	22.57	21.53	**	-	-	-
15	OVARC1001051	381.08	484.89	350.06	354.62	546.22	323.29	741.7	383.07	703.22				
	OVARC1001054	6.6	5.59	5.17	3.06	5.01	2.19	3.89	3.27	2.79	**	-	-	-
	OVARC1001055	14.4	14.98	14.95	7.9	10.31	6.02	5.76	7.98	6.8	**	**	-	-
	OVARC1001062	17.06	13.11	19.77	9.47	7.02	5.92	5.47	7.54	3.76	*	**	-	-
	OVARC1001065	58.8	47.67	46.94	41.39	53.25	31.45	28.27	24.21	23.74	**	-	-	-
	OVARC1001068	12.24	9.85	10	9.05	10.45	10.03	4.46	8.42	5.6	*	-	-	-
20	OVARC1001072	5.2	5.85	5.86	2.66	7.45	1.74	3.96	4.57	2.54	*	-	-	-
	OVARC1001073	7.5	8.58	12.65	4.78	6.4	4.65	5.22	6.96	5.47				
	OVARC1001074	4.7	5.22	6.88	3.02	6.61	3.48	7.1	3.06	2.4				
	OVARC1001078	12.05	11.95	9	4.67	9.88	3.56	7.61	7.71	7.4	*	-	-	-
	OVARC1001085	11.18	10.66	13.33	5.34	5.76	6.26	7.62	7.57	5.14	**	*	-	-
	OVARC1001086	9.39	8.11	10.75	4.2	10	3.47	6.76	8.44	4.74				
25	OVARC1001091	241.22	186.8	193.81	163.03	271.86	178.86	140.22	106.42	112.84	*	-	-	-
	OVARC1001092	18.15	15.63	18.59	22.78	28.04	21.17	13.63	16.45	14.87	*	+	-	-
	OVARC1001104	8.1	6.77	10.2	3.75	7.44	6.44	3.73	7.33	6.12				
	OVARC1001107	107.49	92.87	112.02	98.9	125.4	90.42	58.37	49.66	59.61	**	-	-	-
	OVARC1001113	4.26	4.3	6.18	3.48	7.31	2.59	2.8	2.95	2.48	*	-	-	-
	OVARC1001117	14	17.53	15.78	9.22	10.42	7.27	18.6	16.88	12.72	**	-	-	-
30	OVARC1001118	28.39	26.32	27.08	13.26	13.63	13.64	14.58	18.84	21.16	**	*	-	-
	OVARC1001125	15.09	9.24	18.28	5.27	6.72	6.56	3.81	6.47	3.35	*	*	-	-
	OVARC1001129	8.66	7.06	12.46	3.03	6.25	3.24	5.49	10.45	5.05				
	OVARC1001132	9.08	13.23	15.1	8.86	15.2	9.07	5.57	13.13	8.52				
	OVARC1001138	155.24	193.79	160.3	115.39	149.38	101.68	122.48	164.58	137.03				
35	OVARC1001141	12.99	10.8	12.94	7	9.03	7.57	7.48	10.75	7.3	**	*	-	-
	OVARC1001154	128.63	107.01	106.33	119.46	158.63	83.66	81.42	78.16	83.39	*	-	-	-
	OVARC1001161	17.68	27.92	21.1	11.55	22.82	12.42	16.84	15.27	15.42				
	OVARC1001162	21.74	16.75	19.82	10.26	11.33	9.72	10.1	12.69	11.14	**	**	-	-
	OVARC1001163	10.91	8.03	10.73	4.37	5.95	4.38	3.95	7.35	4.62	**	*	-	-
	OVARC1001167	38.47	38	43.81	17.29	22.87	18.91	28.74	29.72	36.08	**	*	-	-
40	OVARC1001169	11.94	7.48	10.4	5.85	13.24	4.72	3.88	5.51	4.65	*	-	-	-
	OVARC1001170	26.22	26.34	31.69	14.46	15.02	12.36	12.5	14.3	11.17	**	**	-	-
	OVARC1001171	143.81	236.48	129.81	114.4	117.18	158.91	77.44	54.36	128.19				
	OVARC1001173	24.91	33.6	36.93	11.93	16.26	13.53	26.95	27.14	25.91	**	-	-	-
	OVARC1001176	209.32	244.43	234.51	205.75	340.26	228.89	167.09	179.74	200.74	*	-	-	-
	OVARC1001180	67.2	61.17	66.56	30.38	32.42	26.76	42.84	47.13	54.33	**	*	-	-
45	OVARC1001183	18.84	14.15	17.06	4.37	15.93	5.9	10.16	39.68	8.56				
	OVARC1001200	14.51	7.89	14.82	6.11	7.72	3.65	4.16	15.56	6.11				
	OVARC1001202	37.46	21.56	29.17	10.06	10.43	13.25	17.78	23.14	18.29	*	-	-	-
	OVARC1001206	6.83	6.97	11.31	3.06	5.96	2.9	2.03	8.52	4.3				
	OVARC1001209	56.31	51.58	60.2	57.91	69.04	54.41	36.32	29.24	28.87	**	-	-	-
	OVARC1001219	14.82	14.11	12.95	3.65	7.48	4.86	4.34	7.98	3.54	**	**	-	-
50	OVARC1001222	17.11	17.29	18.38	6.61	16.42	9.64	6.56	7.21	5.56	**	**	-	-
	OVARC1001232	34.23	34.49	36.17	16.43	23.42	19.3	24.37	27.16	28.45	**	**	-	-
	OVARC1001240	22.68	22.79	23.45	11.83	13.49	10.07	14	20.54	14.23	**	*	-	-
	OVARC1001243	8.41	5.82	7.86	3.1	7.95	3.99	3.48	18.25	1.41				
	OVARC1001244	43.29	22.42	33.3	20.02	23.25	15.87	15.69	21.46	15.18				
	OVARC1001246	191.8	262.18	216.69	139.76	292.92	181.43	261.49	147.46	197.54				
55	OVARC1001247	33.18	25.43	26.03	8.87	21.81	17.23	8.61	11	7	**	-	-	-
	OVARC1001260	10.58	9.81	13.5	4.35	12.46	3.56	7.06	8.38	13.07				
	OVARC1001261	10.86	9.17	11.3	5.17	7	3.2	4.94	4.19	3.51	*	**	-	-

Table 462

	OVARC1000679	17.17	15.32	16.17	8.84	11.11	9.5	8.07	11.07	12.27	**	*	-	-
	OVARC1000681	8.2	7.24	6.7	4.69	3.61	5.03	3.58	6.1	3.97	**	*	-	-
5	OVARC1000682	113.12	97.53	93.09	104.53	135.81	99.85	59.24	48.57	73.95	*	*	-	-
	OVARC1000689	57.01	42.21	46.96	44.44	54.09	47.18	30.34	27.71	26.26	*	*	-	-
	OVARC1000700	15.99	15.69	20.71	10.16	14.93	8.67	10.07	14.65	13.84	*	*	-	-
	OVARC1000703	26.78	29.43	35.22	14.83	20.62	12.69	19.45	19.63	21.91	*	*	-	-
	OVARC1000722	101.86	85.68	116.52	89.03	99.63	84.39	90.99	59.44	79.26	*	*	-	-
	OVARC1000726	14.33	9.39	12.54	8.8	8.89	8.63	5.72	6.56	7.04	*	*	-	-
10	OVARC1000727	10.18	7.75	13.09	5.79	7.58	4.18	6.38	6.81	6.67	*	*	-	-
	OVARC1000730	29.15	29.95	31.68	12.76	10.82	12.92	11.85	21.92	19.46	**	*	-	-
	OVARC1000741	29.33	29.17	34.55	13.03	37.4	9.12	12.19	21.98	17.49	*	*	-	-
	OVARC1000746	7.78	5.19	7.05	4.39	4.32	3.45	2.05	2.62	3.13	*	**	-	-
	OVARC1000764	38.05	40.13	44.9	31.76	33.36	24.43	25.99	25.42	30.04	*	**	-	-
	OVARC1000769	15.32	19.03	26.3	6.54	8.47	6.17	10.92	9.44	11.01	*	*	-	-
15	OVARC1000771	12.14	10.36	12.36	5.65	9.59	2.99	5.56	6.75	5.39	**	*	-	-
	OVARC1000773	208.54	178.97	168.4	156.38	229.88	197.98	120.99	98.51	115.59	**	*	-	-
	OVARC1000775	34.96	28.91	27.77	13.91	16.38	12.88	14.63	18.43	17.02	**	**	-	-
	OVARC1000778	16.25	14.26	17.29	7.71	11.77	9.22	9.21	13.13	13.25	*	*	-	-
	OVARC1000779	5.51	4.08	7.92	1.14	3.76	1.42	1.72	21.49	16.55	*	*	-	-
20	OVARC1000781	14.72	11.11	14.65	7.53	10.61	4.88	3.59	7.19	4.56	*	**	-	-
	OVARC1000787	16.08	11.69	22.35	5.66	7.21	5.45	9	11.11	9.91	*	*	-	-
	OVARC1000789	23.78	29.78	21.78	12.72	20.51	12.99	10.1	11.61	10.12	**	*	-	-
	OVARC1000800	37.24	52.19	47.92	15.44	23.62	15.47	18.1	25.22	22.47	**	**	-	-
	OVARC1000802	10.69	12.79	19.22	5.61	8.97	4.92	12.57	10.26	7.88	*	*	-	-
	OVARC1000810	41.47	36.61	45.9	14.82	21.04	14.45	26.69	32.64	43.34	**	*	-	-
25	OVARC1000811	19.55	15.32	20.3	9.11	11.03	10.04	5.82	8.76	8.16	**	**	-	-
	OVARC1000814	47.61	40.61	47.27	21.78	30.43	20.51	28.06	40.57	35.2	**	*	-	-
	OVARC1000816	30.44	36.85	45.82	25.26	35.06	20.13	15.73	10.27	23.79	*	*	-	-
	OVARC1000817	7.44	5.99	13.01	2.4	3.28	3.96	1.6	3.36	1.57	*	*	-	-
	OVARC1000834	9.2	11.08	10.6	3.33	5.92	2.58	4.12	4.4	2.95	**	**	-	-
	OVARC1000846	33.2	36.76	34.02	15.54	20.84	16.18	25.4	21.7	22.13	**	**	-	-
30	OVARC1000850	10.1	9.46	13.94	3.02	6.63	4.93	5.97	6.78	5.46	*	*	-	-
	OVARC1000853	124.13	83.82	117.1	50.7	94.43	66.32	58.07	66.46	64.96	*	*	-	-
	OVARC1000862	9.56	7.61	8.86	6.22	4.52	5.06	2.49	2.85	3.78	*	**	-	-
	OVARC1000873	22.12	9.86	11.25	3.95	10.2	7.01	4.33	4	5.56	*	*	-	-
	OVARC1000875	25.32	18.88	20.49	9.43	11.13	12.13	10.43	9.71	12.19	**	**	-	-
	OVARC1000876	14.43	11.76	13.8	4.33	8.64	6.48	3.87	5.38	3.7	**	**	-	-
35	OVARC1000883	16.65	14.83	22.63	10.71	10.46	10.25	4.07	3.52	3.44	*	**	-	-
	OVARC1000885	14.94	15.72	15.2	7.47	13.49	6.08	8.84	6.87	4.31	**	*	-	-
	OVARC1000886	13.43	12.39	14.55	7.87	12.43	7.38	15.33	6.54	7.36	*	*	-	-
	OVARC1000890	238.12	222.49	244.97	168.11	248.06	165.63	120.4	102.6	108.39	**	*	-	-
40	OVARC1000891	14.84	13	11.09	4.07	5.36	4.55	3.08	4.38	3.01	**	**	-	-
	OVARC1000897	5.52	4.78	5.55	2.64	2.48	2.49	2.14	2.77	2.38	**	**	-	-
	OVARC1000912	18.82	12.83	16.34	8.16	8.42	8.22	8.21	8.74	10.52	*	*	-	-
	OVARC1000914	14.59	8.15	9.14	3.85	5.26	5.95	4.68	5.3	4.1	*	*	-	-
	OVARC1000915	17.53	15.34	14.44	6.42	7.74	4.39	7.85	11.07	8.04	**	**	-	-
	OVARC1000916	15.19	18.72	21.06	9.85	20.11	11.22	12.02	12.35	10.4	*	*	-	-
	OVARC1000924	11.12	12.86	14.01	5.55	9.51	4.44	3.13	4.61	3.65	*	**	-	-
	OVARC1000928	46.27	37.28	34.31	5.2	8.81	6.65	4.93	6.15	5.85	**	**	-	-
45	OVARC1000936	10.09	10.55	9.33	5.2	9.07	4.74	3.02	5.31	5.15	**	*	-	-
	OVARC1000937	7.99	8.04	6.58	4.25	5.19	2.94	3.85	3.95	5.45	*	*	-	-
	OVARC1000945	15	14.98	13.39	5.71	5.62	7.78	3.1	4.11	2.76	**	**	-	-
	OVARC1000948	7.88	5.06	4.83	2.65	3.28	2.79	2.69	2.39	2.47	*	*	-	-
	OVARC1000956	14.88	11.07	13.93	5.15	8.05	6.51	8.38	4.46	6.46	**	*	-	-
50	OVARC1000959	16.85	16.96	14.18	9.84	10.41	6.42	15.3	9.78	14.01	**	*	-	-
	OVARC1000960	47.32	52.37	41.49	22.16	32.89	19.82	46.28	34.15	44.56	*	*	-	-
	OVARC1000964	306.84	272.77	304.89	274.01	264.01	211.54	160.52	108.2	102.39	**	*	-	-
	OVARC1000971	7.74	6.08	3.76	3.85	7.47	4.63	2.45	4.82	2.32	*	*	-	-
	OVARC1000975	109.05	95.15	91.82	89.45	126.14	83.84	54.44	48.09	54.61	**	*	-	-
	OVARC1000976	6.41	7.34	5.63	3.4	4.98	2.89	5.48	3.56	2.85	*	*	-	-
55	OVARC1000981	24.82	24.19	24.61	14.13	15.11	11.79	18.79	11.54	16.77	**	*	-	-
	OVARC1000982	21.37	12.18	12.3	7.66	16.73	9.68	5.67	4.54	5.45	*	*	-	-
	OVARC1000984	10.44	9.85	10.17	2.64	4.79	2.24	2.88	2.32	3.54	**	**	-	-
	OVARC1000995	27.57	24.28	19.39	14.42	17.35	10.4	23.31	17.65	22.26	*	*	-	-

Table 461

	OVARC1000313	31.08	25.28	22.97	16.65	33.34	16.32	7.95	14.64	16.92	*	-
	OVARC1000321	95.57	70.17	108.42	49.21	78.85	36.91	71.72	84.76	66.41	*	-
5	OVARC1000326	9.36	7.55	9.19	5.69	4.73	3.83	2.8	4.95	2.81	**	-
	OVARC1000327	9.64	6.54	7.7	4.58	5.69	4.67	3.24	3.34	3.39	*	-
	OVARC1000331	31.51	26.56	17.69	20.2	32.18	25.32	7.77	12.73	9.94	*	-
	OVARC1000335	10.5	10.92	9.98	3.56	6.12	5.55	3.62	2.51	3.12	**	-
	OVARC1000347	22.28	19.59	19.77	14.38	17.51	15.31	11.64	10.94	10.7	*	-
	OVARC1000348	9.03	9.66	11.56	3.27	3.92	2.72	2.5	3.43	2.51	**	-
10	OVARC1000363	17.67	18.03	15.99	8.09	14.09	9.86	7.93	4.93	7.38	*	-
	OVARC1000377	9.05	5.89	9.25	3.23	6.62	3.71	2.88	2.47	4.83	*	-
	OVARC1000382	11.38	10.29	10.44	4.87	5.92	3.1	5	4.78	5.93	**	-
	OVARC1000384	15.04	16.77	12.96	3.9	4.19	3.88	3.11	4.29	3.9	**	-
	OVARC1000401	12.88	8.81	11.8	7.36	8.78	6.88	5.26	4.94	4.67	**	-
	OVARC1000406	452.57	396.91	414.58	242.32	241.63	222.43	202.53	242.65	184.03	**	-
15	OVARC1000407	11.18	13.52	12.76	4.82	6.15	3.43	3.77	3.81	3.74	**	-
	OVARC1000408	143.25	152.71	130.24	110.8	163.58	95.03	87.98	95.98	86.83	**	-
	OVARC1000410	12.11	11.07	15.74	6.71	11.27	6.87	6.11	5.6	3.42	**	-
	OVARC1000411	11.28	8.14	7.71	4.73	4.82	3.45	4.09	4.41	4.68	*	-
	OVARC1000414	15.72	17.28	11.48	7.59	8.9	10.02	14.36	14.47	14.88	*	-
	OVARC1000420	11.3	11.1	10.67	5.43	7.49	5.25	8.15	6.28	6.64	**	-
20	OVARC1000421	21.51	18.12	14.96	8.46	10.6	6.1	8.22	6.83	7.06	*	-
	OVARC1000427	255.84	237.66	229.09	230.15	286.16	201.09	155.75	128.13	139.78	**	-
	OVARC1000431	86.97	107.76	104.05	53.83	76.2	48.18	31.1	18.06	18.56	*	-
	OVARC1000437	18.7	18.85	16.29	10.2	15.26	9.01	9.96	5.69	8.45	*	-
	OVARC1000439	14.55	21.25	13.48	9.92	17.37	9.21	5.86	5.66	4.06	*	-
25	OVARC1000440	37.93	36.45	33.24	14.99	19.86	14.94	10.14	8.93	10.09	**	-
	OVARC1000442	47.65	39.77	34.86	17.92	25.44	19.14	36.52	26.4	37.91	*	-
	OVARC1000443	9.63	12.52	9.02	8.56	33.14	5.68	16.38	5.13	9.47	*	-
	OVARC1000461	8.43	5.22	4.46	4.04	5.64	3.64	3.41	4.7	3.49	*	-
	OVARC1000465	13.05	12.4	8.82	5.16	5.82	3.89	4.59	3.18	2.41	*	-
	OVARC1000466	22.78	19.36	18.72	10.66	17.86	12.45	11.65	7.24	8.12	**	-
30	OVARC1000467	8.03	5.77	8.99	2.45	3.8	2.2	3.29	2.29	3.38	*	-
	OVARC1000470	15.96	13.65	10.41	7.96	9.91	7.26	9.53	10.16	10.25	*	-
	OVARC1000473	12.49	10.05	10.69	6.2	7.49	6.37	10.6	7.2	9.15	**	-
	OVARC1000479	24.56	30.78	25.75	8.77	13.76	7.96	8.07	9.81	9.07	**	-
	OVARC1000484	41.23	29.61	25.62	15.32	17.14	13.32	29.7	26.23	34.72	*	-
	OVARC1000486	15.8	17.22	17.4	11.06	13.85	9.12	14.13	11.53	12.18	*	-
35	OVARC1000486	6.74	8.5	5.88	2.96	8.22	5.46	2.84	3.93	3.9	*	-
	OVARC1000520	8.86	5.17	6.53	2.16	5.01	3.05	4.06	3.97	5.92	*	-
	OVARC1000522	35.51	29.95	29.3	26.12	30.21	22.71	22.42	16.55	18.06	**	-
	OVARC1000526	25.75	25.79	18.89	12.38	14.14	8.6	32.78	19.39	27.42	**	-
	OVARC1000529	17.7	16.61	13.49	7.72	9.26	6.61	12.03	7.5	12.39	**	-
	OVARC1000533	31.08	27.41	22.88	25.25	24.54	12.97	18.02	13.21	18.14	*	-
40	OVARC1000543	11.04	5.87	5.45	4.07	4.48	2.25	3.68	3.22	4.84	*	-
	OVARC1000550	9.45	7.02	5.98	3.29	4.46	2.66	4.35	2.47	4.43	*	-
	OVARC1000553	34.95	36.44	34.04	16.33	24.77	21.97	31.22	31.95	33.2	**	-
	OVARC1000556	22.37	23.81	26.15	14.72	11.54	9.38	19.99	16.63	22.58	**	-
	OVARC1000557	11.61	9.78	7.49	2.51	3.89	2.24	6.45	4.03	5.39	**	-
	OVARC1000561	67.36	70.57	52.24	30.14	35.45	26.39	87.82	47.67	77.14	**	-
45	OVARC1000564	79.74	89.59	94.8	35.3	37.73	54.01	49.59	52.84	37.26	**	-
	OVARC1000573	11.41	12.83	14.29	6.97	10.13	6.51	5.93	10.17	8.55	*	-
	OVARC1000576	235.06	210.51	199.48	203.8	258.47	200.46	142.26	126.19	115.56	**	-
	OVARC1000578	15.29	13.44	14.45	8.15	12.93	8.56	9.81	12.76	11.9	*	-
	OVARC1000581	5.46	3.63	4.65	3.12	3.25	1.85	1.84	1.64	2.35	**	-
	OVARC1000586	99.11	90.08	96.7	58.86	53.54	48.9	78.17	82.33	62.01	**	-
50	OVARC1000588	5.04	6.16	5.9	2.73	4.3	3.79	4.96	4.69	3.2	*	-
	OVARC1000605	11.72	9.54	8.96	6.4	6.63	6.02	4.7	4.72	5.55	*	-
	OVARC1000622	86.95	79.96	78.79	29.56	66.79	40.9	71.63	93.74	75.18	*	-
	OVARC1000636	9.18	7.82	11.99	5.56	6.68	4.67	5.37	6.74	3.7	*	-
	OVARC1000640	9.32	12.02	12.8	8.63	10.33	8.57	6.08	8.89	7.4	*	-
	OVARC1000649	137.37	159.82	123.65	116.33	170.6	128.38	91.38	85.12	81.65	**	-
55	OVARC1000661	14.89	15.01	20.86	11.94	15.89	11.61	9.31	9.27	13.9	*	-
	OVARC1000677	42.34	38.62	28.23	30.67	34.02	28.61	19.25	22.35	28.93	*	-
	OVARC1000678	18.66	21.84	18.64	9.55	15.51	8.53	9.3	11.98	8.4	*	-

Table 460

	WT2RP6000182	11.67	12.98	10.81	11.27	4.67	4.19	7.63	8.13	8.65	**	-
	OVARC1000001	10.18	10.98	7.82	10.48	3.67	5.27	8.05	6.3	4.32	*	+
5	OVARC1000003	19.46	16.04	21.53	24.56	18.39	28.35	24.31	27.93	25.27	*	+
	OVARC1000004	154.65	161.68	111.37	257.56	204.62	269.89	148.7	142.43	153.8	*	+
	OVARC1000006	8.96	9.17	13.23	12.36	9.89	7.09	8.46	11.94	16.91	*	-
	OVARC1000013	12.84	15.42	21.83	13.5	7.52	11.75	8.01	7.31	7.45	*	-
	OVARC1000014	9.51	5.51	9.75	8.67	3	2.57	4.2	4.5	5.23	*	-
	OVARC1000017	7.8	10.89	11.1	9.23	5.72	5.56	7.08	8.83	7.59	*	-
10	OVARC1000026	38.97	98.56	58.17	35.12	21.04	21.52	23	21.16	19.52	*	-
	OVARC1000035	53.62	63.74	56.85	49.18	30.89	52.6	44.01	25.83	35.4	*	-
	OVARC1000037	44.18	27.1	33.64	21.27	17.7	36.96	22.76	28.28	23.56	*	-
	OVARC1000058	21.2	16.03	18.93	19.43	6.93	18.25	18.85	19.08	20.93	*	-
	OVARC1000060	8.69	7.53	9.42	11.74	4.69	15.1	7.86	12.81	10.28	*	-
	OVARC1000068	11.63	6.29	9.06	11.58	8.06	9.18	6.71	8.97	7.36	*	-
15	OVARC1000069	130.28	91.82	110.55	138.44	48.02	81.99	34.87	114.87	91.52	*	-
	OVARC1000071	4.7	2.59	5.56	6.14	1.28	3.78	7.5	3.79	3.09	*	-
	OVARC1000075	608.15	715.16	979.97	596.89	401.8	521.41	1505.9	1295.8	1454	**	+
	OVARC1000083	31.41	22.53	32.47	28.89	10.2	12.73	23.7	17.6	16.65	*	-
	OVARC1000085	150.84	132.35	115.14	65.66	79.36	104.26	57.87	55.95	46.39	*	-
20	OVARC1000086	21.41	22.84	29.16	11.19	16.67	9.71	14.87	19.23	16.21	*	-
	OVARC1000087	5.75	4.1	5.37	4.05	5.46	2.99	2.78	3.2	3.4	*	-
	OVARC1000090	32.43	34.99	39.46	22.05	34.29	19.79	30.39	42.7	35.58	*	-
	OVARC1000091	19.21	16.58	18.4	9.39	12.78	9.69	3.97	5.82	3.09	**	-
	OVARC1000092	12.38	12.28	20.09	9.9	9.89	8.91	11.14	12.55	9.97	**	-
	OVARC1000105	143.72	148.63	164.07	88.33	70.71	97.62	141.54	139.36	177.6	**	-
	OVARC1000106	56.13	50.31	43.64	33.71	26.64	28.64	25.97	22.43	26.91	**	-
25	OVARC1000109	41.67	37.23	42.5	18.44	22.06	21.86	20.05	26.41	16.1	**	-
	OVARC1000113	24.72	16.19	25.29	12.07	22.86	8.51	9.56	13.5	7.9	*	-
	OVARC1000114	20.78	17.23	18.89	9.6	15.6	9.38	19.16	20.46	17.77	*	-
	OVARC1000133	17.57	29.71	58.68	7.88	21.62	13.9	11.06	7.62	51.45	*	-
	OVARC1000137	26.29	21.24	32.71	19.41	23.84	21.06	17.64	15.76	21.29	*	-
	OVARC1000139	159.47	147.34	170.25	146.09	161.46	121.94	114.2	86.17	91.75	**	-
30	OVARC1000145	5.46	5.11	5.9	4.05	7.97	2.4	2.98	4.01	3.61	**	-
	OVARC1000148	26.75	26.35	26.82	10.67	7.97	11.29	11.04	13.8	10.7	**	-
	OVARC1000151	10.86	7.69	7.49	4.61	6.76	7.44	5.23	9.46	5.82	*	-
	OVARC1000157	131.87	72.71	159.16	28.64	46	29.09	32.55	30.3	35.03	*	-
	OVARC1000162	5.48	6.74	10.66	5.09	8.45	3.6	3.17	7.57	4.35	*	-
35	OVARC1000168	20.5	23.24	26.19	11.56	15.79	12.79	11.3	16.45	13.91	**	-
	OVARC1000169	67.7	69.5	69.91	59.55	32.42	51.67	35.89	25.97	33.69	**	-
	OVARC1000178	7.01	7.74	9.93	3.35	8.16	5.02	5.67	3.4	4.49	*	-
	OVARC1000182	6.26	2.63	4.45	2.72	4.87	3.13	2.7	3.24	3.81	*	-
	OVARC1000186	14.26	11.19	12.68	6.9	8.25	3.97	5.99	5.56	7.72	*	-
	OVARC1000188	19.42	11.08	15.15	6.26	7.14	4.44	6.66	6.47	9.02	*	-
40	OVARC1000191	10.3	6.61	7.55	5.45	7.49	3.47	4.08	6.65	4.21	*	-
	OVARC1000198	22.01	20.32	21.51	13.28	15.24	8.22	13.24	17.04	19.28	*	-
	OVARC1000208	131.85	125.05	141.2	106.48	130.32	109.42	107.54	141.64	112.43	*	-
	OVARC1000209	37.5	42.42	46.9	35.76	48.15	33.79	26.69	26.98	20.75	**	-
	OVARC1000212	14.17	16.96	15.84	10.34	14.11	8.11	6.32	5.07	5.47	**	-
	OVARC1000216	7.5	5.29	9.53	2.31	4.11	1.95	4.22	3.32	1.67	*	-
45	OVARC1000240	30.72	25.44	20.48	13.87	16.01	11.11	15.84	21.45	17.35	*	-
	OVARC1000241	8.83	9.47	8.6	4.52	5.75	4.08	4.87	6.55	6.44	**	-
	OVARC1000249	12.5	10.38	14.17	6.99	8.18	4.94	7.53	30.07	10.99	*	-
	OVARC1000254	211.04	164.08	171.72	162.84	218.18	173	104.59	109.93	114.49	**	-
	OVARC1000255	8.62	6.5	15.01	2.8	3.3	2.22	1.58	8.88	3.88	*	-
	OVARC1000267	41.48	57.68	54.05	22.69	31.39	23.55	19.25	29.77	21.42	*	-
50	OVARC1000275	6.58	9.72	13.14	1.2	3.21	1.15	3.78	4.58	2.47	*	-
	OVARC1000287	20.12	19.59	19.09	8.41	13.5	9.21	9.7	16.66	9.51	**	-
	OVARC1000288	34.41	32.21	34.44	10.93	20.1	12.59	9.34	11.6	15.85	**	-
	OVARC1000298	37.89	35.05	40.4	26.93	41.61	26.19	16.03	19.69	20.76	**	-
	OVARC1000302	10.27	8.52	8.63	3.64	6.33	2.17	4.38	8.26	8.64	*	-
	OVARC1000304	13.52	8.67	11.83	6.89	9.55	6.81	7.04	9.75	5.89	*	-
55	OVARC1000307	9.2	9.65	13.33	4.99	7.9	7.07	7.34	4.85	4.2	*	-
	OVARC1000309	10.77	7.81	10.14	3.65	4.52	3.3	3.45	4.14	3.51	**	-
	OVARC1000312	38.52	63.36	48.86	16.29	39.99	22.16	11.44	23.2	11.33	*	-

Table 459

	NT2RP4002377	39.07	34.91	52.74	47.19	54.21	38.25	54.01	36.88	39.01			
	NT2RP4002408	8.23	10.17	7.33	8.55	4.11	5.54	5.29	2.67	3.53	*		-
5	NT2RP4002425	16.72	9.87	22.95	18.36	10.72	16.61	14.74	22.3	29.56			
	NT2RP4002432	134.52	97.55	162.62	149.32	142.05	170.39	110.43	102.21	77.71			
	NT2RP4002447	15.99	16.51	16.61	15.14	7.55	8.09	7.58	13.4	16.29			
	NT2RP4002451	15.11	13.46	19.12	12.6	10.1	7.99	7.15	14.07	11.51			
	NT2RP4002461	94.61	108.24	116.21	78.57	91.21	74.75	80.34	65.93	74.08	*	*	-
	NT2RP4002486	16.89	12.32	16.5	14.11	11.37	5.17	6.32	4.06	4.53	**		-
10	NT2RP4002517	13.38	14.55	13.22	12.72	9.03	11.46	10.8	6.18	8.61	*		-
	NT2RP4002556	14.42	18.7	14.04	18.8	13.59	13.31	16.09	6.06	5.52			
	NT2RP4002569	16.26	11.41	27.18	13.33	6.98	10.11	3.13	6.78	11.56			
	NT2RP4002587	4.28	5.9	13.49	5.9	21.35	3.55	2.37	3.48	4.03			
	NT2RP4002591	11.21	8.73	16.61	10.63	3.25	5.39	2.62	6.98	4.42	*		-
	NT2RP4002607	7.19	4.54	8.62	7.82	4.54	6.78	3.93	3.64	3.85			
15	NT2RP4002627	129.04	84.31	131.91	123.08	121.1	113.29	100.69	103.33	117.09			
	NT2RP4002628	23.42	19.42	25.54	15.3	10.07	9.36	10.28	14.93	5.74	*	*	-
	NT2RP4002630	34.52	45.14	38.61	24.65	19.47	21.97	38.2	24.12	21.41	**		-
	NT2RP4002639	69.58	58.92	77.5	92.97	83.99	82.67	76.62	44.55	74.09	*	+	
	NT2RP4002641	10.87	7.59	20.93	9.6	5.64	6.42	5.77	9.03	8.9			
20	NT2RP4002658	167.82	202.75	171.29	210.71	275.39	155.29	113.71	313.1	177.93			
	NT2RP4002669	11.89	14.59	18.31	13.81	8.54	6.37	6.6	13.8	11.32			
	NT2RP4002677	38.73	39.84	49.87	21.84	19.97	14	8.4	12.92	7.79	**	**	-
	NT2RP4002715	60.14	28.66	66.85	78.28	49.48	49.64	41.22	48.79	51.15			
	NT2RP4002750	5.11	4.54	6.02	5.02	2.86	2.29	1.61	2.5	1.93	**		-
	NT2RP4002784	13.9	15.39	21.97	16.43	16.29	10.56	10.51	11.54	8.19			
	NT2RP4002791	19.14	16.24	15.43	15.81	11.98	5.79	6.28	6.12	6.25	**		-
25	NT2RP4002811	7.07	4.03	15.62	8.02	4.02	6.88	1.73	4.87	5.66			
	NT2RP4002830	22.24	23	26.1	14.14	12.81	9.89	5.49	12.55	7.49	**	**	-
	NT2RP4002832	6.62	7.2	10.83	9.33	4.27	5.54	7.23	5.78	6.45			
	NT2RP4002850	120.8	112.69	112.51	87.95	73.32	75.57	95.07	111.13	108.7	**		-
	NT2RP4002874	8.25	8.24	13.89	8.67	2.89	3.54	1.21	3.45	2.95	*		-
30	NT2RP4002884	89.55	83.99	93.99	87.7	70.28	83.78	63.11	54.66	68.06	**		-
	NT2RP4002888	6.66	8.16	10.03	7.45	4.69	3.72	4.86	3.64	2.94	*		-
	NT2RP4002891	19	20.18	21.15	16.8	8.39	11.91	12.08	8.99	11.53	*	**	-
	NT2RP4002894	39.86	38.92	34.99	21.18	29.78	22.86	11.88	16.42	39.6	*		-
	NT2RP4002896	7.81	11.66	17.8	10.13	3.4	6.5	5.01	4.37	5.08			
	NT2RP4002905	5.89	3.63	14.35	7.98	3.09	2.98	3	5.21	1.93			
35	NT2RP4002907	14.82	12.9	19.78	10.14	7.04	6.74	11.86	12.75	9.47	*		-
	NT2RP5003459	437.62	468.29	451.9	449.8	544.78	414.46	466.17	464	478.98			
	NT2RP5003461	21.93	17.54	18.05	12	9.18	10.65	7.44	6.63	4.58	**	**	-
	NT2RP5003471	52.72	62.97	42.21	31.11	21.1	25.12	46.83	20.37	35.04	*		-
	NT2RP5003477	5.7	3.4	3.74	5.93	3.44	2.66	3.17	2.23	2.28			
	NT2RP5003487	1749.6	1957.9	1778.8	1241.4	1573.1	1807.7	1562.4	2343.9	2243.6			
40	NT2RP5003492	6.08	6.5	15.88	6.67	3.98	3.96	2.59	3.25	1.92			
	NT2RP5003500	4.26	2.52	9.75	6.65	2.34	2.66	2.29	2.53	1.57			
	NT2RP5003506	10.49	9.24	16.38	12.74	8.52	9.22	9.69	9.29	9.27			
	NT2RP5003512	6.99	7.06	8.63	5.95	2.46	3.36	2.76	2.88	3.49	*	**	-
	NT2RP5003522	10.07	9.26	8.55	5.18	3.12	3	5.35	1.85	3.07	**	**	-
	NT2RP5003524	6.44	9.05	5.63	7.13	1.41	1.72	4.66	2.23	3	*		-
45	NT2RP5003527	117.46	141.32	115.3	136.41	95.19	117.09	134.61	57.44	67			
	NT2RP5003531	24.59	24.67	34.31	22.85	17.67	26.56	12.06	25.82	35.99			
	NT2RP5003534	6.96	7.88	13.1	6.91	5.36	2.92	2.47	4.02	3.38	*		-
	NT2RP6000020	19.43	27.72	47.01	19.42	30.65	12.48	29.24	22.48	15.27			
	NT2RP6000022	13.58	7.81	13.25	8.64	4.5	5.83	7.1	5.5	4.05	*		-
	NT2RP6000050	5.27	8.58	7.67	6.81	3.59	4.26	5.19	3.06	5.18			
50	NT2RP6000063	5.9	4.93	5.56	6.03	1.07	1.6	2.56	2.13	3.15	**		-
	NT2RP6000074	10.12	8.88	7.2	8.89	3.35	3.32	4.76	3.03	4.96	*		-
	NT2RP6000083	36.48	50.53	36.06	34.94	19.9	25.84	29	16.57	36.99			
	NT2RP6000100	9.33	8.85	11.79	8.7	3.93	5.47	3.33	5.54	5.88	*		-
	NT2RP6000123	7.59	5.31	8.35	10.56	5.66	4.44	3.17	5.91	3.98			
	NT2RP6000129	4.7	6.51	7.04	5.86	2.6	3.34	4.3	7.17	3.53			
55	NT2RP6000147	23.98	19.21	18.37	28.9	18.17	21.68	18.73	17.92	16.85			
	NT2RP6000163	6.02	5.5	6.17	6.66	2.33	2.81	2.76	2.24	3.12	**		-
	NT2RP6000181	31.82	37.36	38.55	11.76	5.01	6.27	13.35	6.28	18.64	**	**	-

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	NT2RP4001927	22.95	20.54	20.03	14.02	11.73	7.04	7.86	6.56	6.03	*	**	-	-
	NT2RP4001931	31.71	30.66	39.53	15.12	11.86	11.1	6.42	8.14	7.84	**	**	-	-
5	NT2RP4001933	124.92	135.82	123.56	117.23	83.52	79.43	61.4	54.06	63.46		**	-	-
	NT2RP4001938	21.23	9.81	28.3	17.33	8.99	14.74	12.02	19.38	17.02			-	-
	NT2RP4001942	40.71	46.99	41.11	32.73	26.67	27.19	19.11	35.23	32.36	**		-	-
	NT2RP4001945	3.45	3.93	9.62	6.08	1.55	2.05	2.15	2.81	1.6			-	-
	NT2RP4001946	10.3	8.04	13.38	10.29	3.79	3.79	3.46	4.08	3.41	*	*	-	-
10	NT2RP4001947	16.45	15.77	21.11	11.34	9.6	7.02	4.68	3.24	4.94	*	**	-	-
	NT2RP4001950	5.04	4.87	6.27	6.87	2.41	1.7	3	2.63	1.3		*	-	-
	NT2RP4001953	26.47	19.18	22.05	22.48	9.35	15.94	17.83	14.03	17.8		*	-	-
	NT2RP4001966	6.74	6.56	3.55	6.86	3.2	2.07	1.97	1.5	1.8		*	-	-
	NT2RP4001970	7.84	6.46	14.19	9.16	3.49	5.37	2.96	4.43	3.21			-	-
	NT2RP4001975	16.36	22.03	28.3	17.31	12	7.09	9.7	16.29	13.08			-	-
15	NT2RP4001988	38	36.95	48.12	29.2	31.09	30.9	21.07	21.03	23.16	*	**	-	-
	NT2RP4001996	24.73	23.55	24.27	15.39	14.66	5.03	8.32	9.1	9.68	*	**	-	-
	NT2RP4002014	11.39	15.95	16.62	10.15	7.55	6.98	8.4	7.61	5.52	*	*	-	-
	NT2RP4002018	16.73	14.11	11.76	12.16	9.31	11.09	15.09	12.65	8.98			-	-
	NT2RP4002035	5.82	5.03	4.44	7.72	3.9	4.37	5.69	2.8	2.78			-	-
	NT2RP4002043	18.95	25.2	14.18	17.22	10	14.87	15.39	7.04	10.89			-	-
20	NT2RP4002046	24.11	28.29	25.84	19.27	12.78	20.33	11.41	17.55	17.78	*	*	-	-
	NT2RP4002052	12	15.65	20.88	9.3	9.08	6.85	4.02	3.61	4.24	*	**	-	-
	NT2RP4002056	153.48	150.03	95.32	64.65	34.92	59.97	51.1	38.55	53.86	*	*	-	-
	NT2RP4002057	40.68	39.2	39.83	19.31	30.56	26.73	27.83	37.55	52.18	*	*	-	-
	NT2RP4002058	17.65	14.46	13.17	10.87	3.28	4.52	5.26	2.89	2.9	*	**	-	-
	NT2RP4002064	5.15	7.64	7.19	5.68	1.88	1.94	2.82	1.16	2.09	**		-	-
25	NT2RP4002071	19.67	23.04	16.87	12.91	7.48	6.56	8.07	4.85	9.75	*	**	-	-
	NT2RP4002075	4.49	2.91	8.1	6.42	3.43	5.27	3.76	2.11	2.28			-	-
	NT2RP4002078	19.28	11.77	21.52	11.04	6.62	7.49	7.2	13.73	14.2	*		-	-
	NT2RP4002081	26.93	23.83	24.19	19.25	16.65	23.73	20.7	21.48	20.63	*	*	-	-
	NT2RP4002083	12.48	9.47	12.44	9.18	6.34	7.57	7.79	10.15	9.98	*		-	-
	NT2RP4002099	4.69	4.27	5.09	6.55	1.89	1.66	2.86	2.04	1.79	**		-	-
30	NT2RP4002106	48.22	85.51	111.5	22.9	29.06	16.21	28.91	8.21	26.28	*	*	-	-
	NT2RP4002111	21.8	28.93	25.95	13.09	13.73	11.32	21.21	10.01	16.62	**		-	-
	NT2RP4002112	18.8	22.05	20.04	17.73	8.26	13.72	15.37	6.12	8.09	*		-	-
	NT2RP4002116	131.42	129.9	145.31	124.89	86.8	130.87	116.59	193.03	130.26			-	-
	NT2RP4002122	9.21	7.05	12.08	10.75	5.68	5.1	3.65	7.4	5.23			-	-
35	NT2RP4002126	16.25	10.45	26.23	9.01	12.04	10.44	10.14	17.51	8.7			-	-
	NT2RP4002133	59.92	73.72	76.92	61.59	43.1	56.16	80.78	84.66	86.01			-	-
	NT2RP4002136	12.24	16.73	29.34	16.04	8.23	7.86	20.93	16.43	17.62			-	-
	NT2RP4002139	65.39	168.51	135.79	98.21	29.79	27.09	86.97	65.15	29.84			-	-
	NT2RP4002174	10.15	13.02	11.19	12.5	4.36	7.32	8.88	11.55	11.53			-	-
	NT2RP4002185	23.38	25.82	19.72	13.23	6.3	10.39	10.9	7.33	11.09	**	**	-	-
40	NT2RP4002186	16.88	7.65	12.65	11.9	7.3	14.74	12.63	14.19	17.64			-	-
	NT2RP4002187	28.59	22.57	36.35	26.74	19.36	12.02	11	23.37	23.58			-	-
	NT2RP4002188	5.08	7.53	8.31	11.66	6.59	7.86	4.55	4.87	6.3			-	-
	NT2RP4002199	5.25	3.74	5.62	11.93	2.82	3.8	2.79	4.05	2.9			-	-
	NT2RP4002206	3.81	1.86	2.83	8.57	1.89	1.73	3.4	4.64	3.09			-	-
	NT2RP4002210	5.45	3.2	5.65	5.42	1.97	2.98	3.71	3.22	3.36			-	-
	NT2RP4002222	6.38	6.14	7.04	6.83	3.57	3.25	6.5	5.55	8.67			-	-
45	NT2RP4002241	19.38	17.71	14.08	18.19	11.3	8.8	22.47	17.63	22.91			-	-
	NT2RP4002248	35.82	21.31	27.47	30.46	20.08	26.03	19.35	24.1	27.06			-	-
	NT2RP4002250	1.58	1.78	7.25	3.8	1.3	0.25	0.61	0.65	1.92			-	-
	NT2RP4002259	5.84	5.18	13.48	7.69	2.79	5.42	4.19	2.88	3.6			-	-
	NT2RP4002268	35.43	25.77	32.22	22.21	30.41	16.71	15.9	32.72	23.26			-	-
50	NT2RP4002288	22.95	20.57	22.28	19.59	16.95	19.16	16.63	15.69	17.98	*	**	-	-
	NT2RP4002290	13.7	13.76	25.96	11.44	11.61	14.66	7.05	8.48	10.18			-	-
	NT2RP4002298	10.23	12.39	16.74	11.6	8.3	6.18	9.56	4.97	3.02			-	-
	NT2RP4002306	15.16	15.97	20.14	18.18	10.48	11.51	13.4	11.06	13.62			-	-
	NT2RP4002308	8.45	5.65	13.02	8.55	4	9.83	3.47	4.84	5.83			-	-
	NT2RP4002336	36.27	19.86	28.94	13.28	11.25	15.46	7.56	12.49	16.8	*	*	-	-
	NT2RP4002340	2.84	2.81	9.7	4.27	0.88	0.87	1.22	1.79	3.04			-	-
55	NT2RP4002361	9.6	7.94	13.29	10.99	5.46	5.57	5.78	8.11	5.48			-	-
	NT2RP4002367	2.79	2.98	7.18	5.28	1.95	2.56	3.24	1.85	1.33			-	-
	NT2RP4002368	24.68	18.92	33.22	37.33	27.57	13.68	19.74	22.15	15.46			-	-

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	NT2RP4001555	5.61	4.66	4.4	6.85	1.41	2.54	5.43	2.64	2.46			
	NT2RP4001567	14.43	18.57	12.18	9.17	4.93	4.07	5.01	3.84	5.68	*	**	-
5	NT2RP4001568	20.92	33.87	32.23	19.47	17.65	14.08	25.04	13.01	8.31			
	NT2RP4001569	38.32	29.49	40	37.57	26.2	39.13	21.18	30.23	30.47			
	NT2RP4001571	15.01	7.71	15.89	14.4	7.44	9.88	7.38	9.18	8.27			
	NT2RP4001574	25.53	25.21	25.8	22.48	16.52	21.02	13.78	20.38	15.43	*	*	-
	NT2RP4001575	5.89	6.11	13.99	11.72	4.86	9.13	5.91	5.72	3.19			
	NT2RP4001578	138.32	156.83	176.17	106.73	98.03	117.94	140.07	117.64	162.41	*		-
10	NT2RP4001582	9.9	23.21	17.14	17.16	3.08	11.08	13.53	10.53	12.39			
	NT2RP4001593	39.07	48.78	46.75	25.68	17.84	15.12	20.23	20.2	20.66	**	**	-
	NT2RP4001605	8.89	11.75	9.28	6.36	2.98	4.4	4.38	3.06	5.09	*	**	-
	NT2RP4001606	18.77	11.14	16.78	12.15	11.21	15.99	4.83	13.12	16.6			
	NT2RP4001607	6.86	5.47	8.59	11.52	4.98	6.05	4.83	4.03	5.51			
	NT2RP4001610	3.61	1.38	4	7.17	1.24	4.9	2.72	3.85	3.06			
15	NT2RP4001614	9.32	7.4	5.36	10.94	2.88	8.46	8.3	5.56	6.1			
	NT2RP4001623	4.1	3.9	7.53	8.2	4.59	6.06	6.02	5.72	5.27			
	NT2RP4001626	25.96	32.39	24.45	30.25	8.64	19.85	8.25	9.34	7.78	**		-
	NT2RP4001634	5.5	5.13	5.73	8.25	3.86	4.21	6.58	3.67	4.85			
	NT2RP4001638	6.66	10.41	10.2	8.73	4.08	5.57	10.07	5.59	6.69			
20	NT2RP4001644	54.14	31.27	53.8	60.84	32.15	47.2	22.38	29.74	30.92			
	NT2RP4001646	19.49	10.73	26.93	21.07	12.28	14.22	11.82	16.28	16.41			
	NT2RP4001656	13.27	8.07	17.13	8.91	6.79	8.31	7.24	7.86	6.43			
	NT2RP4001666	5.12	4.37	7.8	6.14	1.29	4.07	3.21	5.08	2.86			
	NT2RP4001670	3.97	4.22	8.36	10.93	2.63	2.89	3.57	4.05	2.82			
	NT2RP4001677	60.48	67.97	86.27	37.88	33.85	40.42	60.13	40.04	66.44	*		-
	NT2RP4001679	63.4	46.03	74.85	69.08	70.96	72.54	80.35	59.3	58.1			
25	NT2RP4001685	25.41	48.09	50.45	43.83	20.99	24.93	40.69	28.74	39.61			
	NT2RP4001696	10.8	7.02	11.48	9.54	3.39	6.61	2.58	6.02	6.23			
	NT2RP4001699	7.34	4.14	12.45	8.78	2.82	5.44	4.48	5.75	8.17			
	NT2RP4001717	14.1	8.06	13.01	8.07	5.02	5.29	6.3	4.01	4.24	*		-
	NT2RP4001719	12.35	7.72	11.2	8.9	5.51	7.17	4.84	7.79	6.21			
	NT2RP4001725	3.63	2.94	5.27	5.13	1.69	2.36	2.09	2.38	2.12			
30	NT2RP4001726	27.43	19.04	36.39	17.49	15.05	17.67	16.15	9	13.24			
	NT2RP4001730	17.07	14.23	28.12	18.8	22.34	22.39	15.23	16.15	19.5			
	NT2RP4001739	13.08	27.52	18.78	28.99	19.24	16.38	17.13	15.51	14.38			
	NT2RP4001741	36.21	33.9	43.76	24.77	17.24	19.76	19.65	43.11	31.46	**		-
	NT2RP4001753	17.2	14.59	16.48	7.48	4.99	6.73	3.91	7.96	13.47	**		-
	NT2RP4001760	8.36	7.54	12.21	7.55	2.73	5.75	3.77	6.5	5.81			
35	NT2RP4001787	449.13	316.48	382.02	579.25	624.67	556.76	483.34	493.96	418.4	**	+	
	NT2RP4001790	7.47	3.89	8.52	8.8	3.39	5.62	4.59	3.58	5.07			
	NT2RP4001795	63.42	46.97	73.29	43.26	32.42	28.05	20.69	20.32	20.73	*	**	-
	NT2RP4001803	8.84	7.88	10.54	9.34	4	5.93	4.96	3.55	3.59	**		-
	NT2RP4001805	5.95	5.69	10.38	8.15	3.79	5.46	9.95	6.17	7.95			
40	NT2RP4001809	55.79	43.66	54.44	43.11	38.36	44.76	37.5	36.54	53.86			
	NT2RP4001817	34.65	23.28	22.77	22.82	15.26	16.99	9.01	16.13	16.73	*		-
	NT2RP4001822	6.52	3.81	10.26	6.23	2.19	2.7	2.67	9.77	2.8			
	NT2RP4001823	3.6	2.46	7.23	4.29	1.65	1.68	0.91	2.7	2.37			
	NT2RP4001827	92.63	82.49	90.1	26.32	25.71	23.6	20.01	17.73	25.06	**	**	-
	NT2RP4001828	61.47	38.75	56.29	48.73	53.93	46.7	40.24	45.94	36.84			
45	NT2RP4001835	44.8	31.89	48.09	43.58	41.3	52.3	39.14	27.95	48.98			
	NT2RP4001838	6.15	7.83	5.55	9.67	6.83	8.27	7.38	14.64	4.59			
	NT2RP4001841	120.68	83.99	187.16	139.43	130.99	191.28	92.44	82.4	82.84			
	NT2RP4001849	5.44	4.67	11.88	5.66	3.18	2.21	2.03	4.71	2.38			
	NT2RP4001861	120.57	86.96	121.65	102.03	95.48	118.7	109.17	140.23	134.58			
	NT2RP4001877	47.77	41.26	29.9	28.86	24.84	22.31	17.4	14.34	21.88			
50	NT2RP4001879	28.67	26.1	20.85	19.88	15.94	14.99	7.59	15.66	4.85	*	*	-
	NT2RP4001889	38.95	24.39	36.19	23.27	20.66	27.73	35.61	41.13	33.21			
	NT2RP4001893	6.77	9.6	11.58	7.5	6.4	5.73	3.91	4.94	4.11	*		-
	NT2RP4001896	20.14	21.68	20.53	23.47	19.99	18.99	15.61	11.88	13.51	**		-
	NT2RP4001898	54.69	54.01	78.1	65.84	57.05	63.77	28.3	45.74	52.57			
	NT2RP4001901	17.19	14.63	24	14.51	10.05	13.25	15.49	18.92	20.72			
55	NT2RP4001910	121.38	105.48	128.54	129.33	152.14	104.65	69.31	116.85	87.82			
	NT2RP4001925	17.35	19.33	26.78	14.34	7.91	7.52	5.99	7.67	6.41	*	**	-
	NT2RP4001926	13.94	13.08	14.09	14.02	5.92	2.48	4.72	6.01	4.66	**		-

Table 456

	NT2RP4001213	7.81	5.05	8.73	8.57	3.73	3.8	4.8	3.72	3.86			
	NT2RP4001214	3.93	5.31	4.9	6.79	4.2	2.68	6.35	2.95	4.56			
5	NT2RP4001219	16.88	22.36	19.11	23.96	11.89	15.23	12.2	5.76	8.23	*		-
	NT2RP4001228	31.68	26.26	28.99	28.25	18.75	24.21	24.62	30.13	29.4			
	NT2RP4001235	19.17	16.01	20.65	14.75	7.88	8.31	4.98	7.75	11.8	*	*	-
	NT2RP4001256	7.87	5.89	15.98	8.52	4.59	3.98	3.22	7.66	6.44			
	NT2RP4001257	10.17	7.29	12.91	9.33	4.43	6.16	4.58	7.66	4.3			
	NT2RP4001260	9	7.35	9.45	7.52	2.82	3.63	1.87	4.5	2.81	**		-
10	NT2RP4001261	14.73	14	19.92	14.97	13.67	11.83	31.07	30.65	37.38	**		+
	NT2RP4001274	46.98	48.98	34.12	11.26	8.82	9.09	7.45	6.66	5.77	**	**	-
	NT2RP4001276	24.77	28.67	28.03	20.15	20.05	19.84	15.05	9.27	8.86	**	**	-
	NT2RP4001283	274.25	154.44	255.86	208.49	145.59	283.17	127.22	121.38	88.24	*		-
	NT2RP4001299	70.57	64.2	62.96	46.39	43.31	37.19	12.59	32.14	14.83	**	**	-
	NT2RP4001313	1.97	3.19	11.08	5.06	1.72	1.96	1.4	3.15	1.49			
15	NT2RP4001315	13.31	15.88	15.77	13.24	9.24	10.09	4.21	7.24	8.5	*	**	-
	NT2RP4001320	57.94	46.23	43.52	51.85	61.21	48.26	36.69	35.49	37.13	*		-
	NT2RP4001325	218.13	214.83	355	324.95	260.73	208.15	240.45	201.7	209.88			
	NT2RP4001336	66.24	37.07	61.11	58.79	61.6	67.7	54.66	37.6	43.21			
	NT2RP4001339	6.71	9.24	9.26	9.34	4.49	4.79	2.82	4.96	4.68	*		-
20	NT2RP4001343	117.66	66.55	122.62	134.91	97.85	136.86	58.06	81.43	87.93			
	NT2RP4001344	93.3	59.49	108.06	106.25	89.38	122.67	65.91	65.77	78.19			
	NT2RP4001345	8.99	7.64	17.74	11.99	4.82	6.81	4.98	6.9	5.71			
	NT2RP4001351	19.43	14.61	27.02	22.21	10.47	17.51	14.3	12.05	12.26			
	NT2RP4001353	3.32	5.15	8.01	5.72	0.75	1.22	1.09	1.47	0.82	*		-
	NT2RP4001355	4.79	4.23	4.91	4.97	2.19	1.13	2.63	2	2.13	**		-
25	NT2RP4001367	14.2	16.94	22.96	11.42	11.61	7.99	7.87	8	4.49	*		-
	NT2RP4001372	4.19	2.77	5.41	7.79	3.55	1.91	2.83	1.62	2.32			
	NT2RP4001373	51.47	33.24	43.9	16.16	26.86	13.77	10.57	24.44	16.81	*	*	-
	NT2RP4001375	4.45	4.59	11.77	7.62	2.36	2.63	2.14	3.81	2.97			
	NT2RP4001379	5.44	4.91	12.41	7.76	2.24	3.26	2.93	5.12	3.63			
	NT2RP4001381	28.02	25.15	21.15	23.38	16.12	13.96	17.03	26.57	24.23			
30	NT2RP4001386	29.66	19.44	36.04	29.55	18.78	18.25	18.75	24.44	26.66			
	NT2RP4001389	11.24	11.58	9.67	9.26	6.9	5.34	3.69	2.72	3.74	*	**	-
	NT2RP4001396	5.21	4.67	8.34	6.72	2.25	1.78	3.47	3.22	1.99			
	NT2RP4001407	5.66	5.49	6.51	7.81	4.54	1.79	3.65	1.8	5.23			
	NT2RP4001409	8.48	7.57	16.74	7.93	2.96	5.53	4.77	6.78	4.46			
35	NT2RP4001410	76.31	89.86	99.62	25.06	52.86	47.85	42.92	82.06	57.7	*		-
	NT2RP4001414	30.84	57.5	63.9	31.08	20.37	24.44	13.71	23.74	21.51	*		-
	NT2RP4001424	11.24	11.55	15.27	11.2	4.99	6.97	6.22	6.82	6.88	**		-
	NT2RP4001433	19.8	14.14	16.87	12.54	11.03	6.56	10.55	8.89	6.43	*	*	-
	NT2RP4001438	69.21	52.84	81.07	58.91	49.01	54.6	39.37	44.33	43.61	*		-
	NT2RP4001442	7.93	5.58	7.34	6.34	2.77	3.09	4.34	2.59	3.24	*		-
	NT2RP4001447	7.95	6.53	5.53	6.92	3.73	3.72	3.38	1.74	2	**		-
40	NT2RP4001466	26.98	30.25	30.96	13.22	11.9	17.04	5.41	8.17	9.81	**	**	-
	NT2RP4001467	56.03	66.43	78.02	42.37	52.35	39.87	17.7	35.75	26.38	*	**	-
	NT2RP4001472	23.78	41.35	30	13.57	13.9	13.93	11.58	12.27	10.38	*	*	-
	NT2RP4001474	7.87	8.46	14.35	7.2	5.31	5.23	4.32	3.7	3.69	*		-
	NT2RP4001483	6.68	4.49	11.62	7.08	3.5	3.8	3.02	2.79	3.04			
45	NT2RP4001488	25.21	54.85	68.99	28.07	36.56	22.06	36.35	46.46	23.9			
	NT2RP4001492	10.33	5.85	8.11	7.33	2	4.79	7.35	5.55	5.24			
	NT2RP4001498	7.69	4.58	4.14	7.14	3.13	3.73	4.3	3.05	2.12			
	NT2RP4001502	209.76	206.24	150.31	118.11	105.88	95.3	44.44	56.95	57.25	*	**	-
	NT2RP4001503	7.44	8.32	13.64	9.66	7.22	6.9	4.97	6.76	6.61			
	NT2RP4001507	13.79	9.83	18.56	10.78	5.48	8.77	10.23	16.21	17.48			
	NT2RP4001510	5.75	3.91	6.87	6.33	3.38	4.41	4.72	4.9	4.63			
50	NT2RP4001516	8.14	6.69	9.41	7.53	2.26	3.8	4.88	3.25	3.25	*		-
	NT2RP4001520	131.31	173.55	237.05	131.12	129.89	116.61	94.53	82.4	103.86			
	NT2RP4001523	33.88	38.97	38.29	18.41	8.08	12.46	21.96	12.77	13.03	**	**	-
	NT2RP4001524	21.05	28.36	29.26	13.82	9.16	8	8.02	5.86	9.68	**	**	-
	NT2RP4001529	19.64	17.51	18.7	21.14	10.37	22.92	12.83	17.42	17.09			
	NT2RP4001531	12.2	10.44	17.59	14.87	8.44	10.35	9.15	12.78	10.56			
55	NT2RP4001546	200.55	317.08	217.26	217.42	252.35	233.58	165.76	290.78	247.68			
	NT2RP4001547	50.85	72.72	70.29	50.48	41.66	51.65	70.8	75.31	63.14			
	NT2RP4001551	5.28	7.23	6.07	6.96	2.03	2.81	2.74	2.79	2.73	**		-

Table 455

	NT2RP4000949	10.71	9.2	16.65	6.86	2.65	3.74	2.78	1.39	2.41	*	*	-	-
	NT2RP4000955	5.81	4.35	6.4	6.28	2.77	1.08	2.31	3.19	3.37	*	*	-	-
5	NT2RP4000959	46.28	46.79	57.49	28.43	49.02	25.11	20.47	30.52	16.93	**	*	-	-
	NT2RP4000962	11.98	9.47	12.39	11.59	10.69	8.83	6.01	4.92	5.77	**	*	-	-
	NT2RP4000973	21.72	16.11	23.28	13.29	14.47	13.18	9.73	13.42	15.34	*	*	-	-
	NT2RP4000975	12.7	11.46	22.76	9.51	5.19	6.19	5.82	10.95	9.25			-	-
	NT2RP4000979	15.15	9.34	21.99	15.27	12.98	9.04	9.28	10.97	8.24			-	-
	NT2RP4000984	6.41	8.08	14.46	6.57	1.97	3.73	2.22	3.31	3.09			-	-
10	NT2RP4000986	6.86	5.41	12.86	6.12	4.31	1.27	1.5	2.5	1.16	*	*	-	-
	NT2RP4000988	17.65	11.88	13.93	10.42	7.43	7.63	9.24	12.28	11.72	*	*	-	-
	NT2RP4000989	4.65	7.43	6.1	6.61	2.52	1.81	2.98	2.09	3.85	*	*	-	-
	NT2RP4000990	6.25	7.92	5.23	9.42	4.63	4.12	5.36	2.51	2.44			-	-
	NT2RP4000994	8.9	9.21	16.92	10.87	4.94	9.69	11.79	23.69	21.99			-	-
	NT2RP4000996	77.3	49.17	79.6	45.61	34.69	39.35	32.71	38.95	46.1	*	*	-	-
15	NT2RP4000997	122.55	129.24	107.05	94.09	70.8	26.94	46.25	80.17	57.14	**	*	-	-
	NT2RP4001001	12.46	18.44	15.8	14.39	8.88	9.17	8.14	8.28	5.33	*	*	-	-
	NT2RP4001004	5.22	3.76	7.06	6.01	1.06	2.64	1.83	1.39	1.05	*	*	-	-
	NT2RP4001006	13.89	13.25	17.25	8.9	7.61	7.07	7.25	6.36	7.3	**	*	-	-
	NT2RP4001009	16.48	20.86	24.07	12.83	13.33	10.38	13	7.19	7.56	*	*	-	-
20	NT2RP4001010	12.07	13.64	9.65	9.47	5.84	7.62	5.8	4.44	3.25	**	*	-	-
	NT2RP4001013	109.49	147.49	90.54	50.6	80.74	63.83	38.95	52.28	51.67	*	*	-	-
	NT2RP4001029	20.54	17.68	29.5	9.58	9.72	9.68	7.27	4.7	8.7	*	*	-	-
	NT2RP4001036	9.27	12.23	13.79	8.52	7.63	4.92	8.03	6.75	4.44	*	*	-	-
	NT2RP4001041	36.4	40.27	32.69	14.35	15.64	11.72	23.48	9.92	16.78	**	*	-	-
	NT2RP4001042	15.67	10.3	17.88	10.18	7.53	8.42	5.86	5.84	7.9	*	*	-	-
25	NT2RP4001046	36.69	45.09	54.4	22.18	26.84	16.1	31.95	24.71	14.49	*	*	-	-
	NT2RP4001050	14.02	22.04	12.83	10.97	2.91	3.84	9.37	3.59	3.94	*	*	-	-
	NT2RP4001051	21.06	26.13	23.41	17.4	11.41	16.25	15.66	11.32	18.06	*	*	-	-
	NT2RP4001057	5.02	3.25	6.98	7.18	2.13	3.82	4.33	2.83	3.14			-	-
	NT2RP4001063	6.13	4.37	7.2	7.29	2.91	4.39	2.87	3.18	3.35	*	*	-	-
	NT2RP4001064	9.3	8.96	15.17	12.02	5.98	5.07	7.19	9.48	7.97			-	-
30	NT2RP4001067	10.58	14.37	13.96	6.93	4.54	4.48	7.26	9.16	8.36	**	*	-	-
	NT2RP4001078	4	5.51	5.6	4.49	1.94	2.06	3.66	2.7	1.04	*	*	-	-
	NT2RP4001079	8.77	14.47	11.97	8.98	3.49	1.98	7.36	2.56	7.44			-	-
	NT2RP4001080	8.56	8.71	5.38	7.24	5.74	5.86	5.43	5.54	6.58			-	-
	NT2RP4001086	8.1	13.85	11.2	8.73	5.59	7.65	7.59	3.27	6.54			-	-
	NT2RP4001095	21.9	10.2	20.98	20.1	8.21	22.42	25.97	28.91	25.72			-	-
35	NT2RP4001098	9.87	7.14	14.73	8.69	4.17	6.42	4.56	7.67	4.16			-	-
	NT2RP4001100	87.64	83.13	87.53	54.87	47.17	71.93	63.14	104.74	100.86	*	*	-	-
	NT2RP4001105	4.62	5.85	6.18	8.6	3.1	3.62	2.72	2.96	3.35	**	*	-	-
	NT2RP4001110	4.35	3.52	3.95	6.4	1.65	2.39	2.86	1.95	2.46	*	*	-	-
	NT2RP4001115	9.9	15.92	16.54	12.73	4.3	5.05	12.1	7.78	11.14			-	-
40	NT2RP4001117	47.61	50.63	46.56	30.24	19.2	17.67	37.4	37.77	12.69	**	*	-	-
	NT2RP4001122	7.49	13.49	11.17	8.93	3.06	4.64	5.55	4.07	4.81	*	*	-	-
	NT2RP4001123	15.99	10.87	15.29	13.96	11.8	8.64	4.92	11.31	11.27			-	-
	NT2RP4001126	32.2	22.75	27.43	23.25	13.48	17.12	8.86	11.87	13.23	**	*	-	-
	NT2RP4001127	5.7	5.54	5.37	7.61	3.72	4.66	3.57	2.2	3.61	**	*	-	-
	NT2RP4001138	6.95	3.88	7.12	9.78	5.56	5.09	4.09	4.44	3.91			-	-
	NT2RP4001143	12.5	11.65	11.43	11.35	7.29	7.76	9.34	8.18	11.45			-	-
45	NT2RP4001148	6.15	6.48	10.09	9.88	3.88	5.87	3.55	2.98	4.94			-	-
	NT2RP4001149	10.56	12.29	14.62	14.76	6.44	4.57	8.21	6.46	7.23	*	*	-	-
	NT2RP4001150	2.08	2.58	2.6	5.53	2.21	2.02	2.86	2.61	5.29			-	-
	NT2RP4001159	28.55	18.66	30.75	17.14	15.33	13.97	7.91	12.94	18.83			-	-
	NT2RP4001162	19.05	7.92	22.71	10.51	6.28	6.36	7.29	6.36	7.28			-	-
	NT2RP4001170	4.82	3.9	7.97	5.23	1.81	3.87	2.53	3.13	2.09			-	-
50	NT2RP4001174	28.66	24.87	30.19	18.62	12.05	22.33	28.65	25.07	34.84	*	*	-	-
	NT2RP4001175	41.57	44.74	48.16	32.54	33.99	41.64	26.8	24.72	38.38	*	*	-	-
	NT2RP4001176	940.52	839.83	1091.9	860.36	609.5	960.82	721.44	858.51	780.83			-	-
	NT2RP4001184	63.86	45.01	63.95	60.3	52	64.4	59.44	58.17	76.19			-	-
	NT2RP4001188	36.14	14.18	33.12	25.99	15.11	24.84	10.23	17.15	15.95			-	-
	NT2RP4001199	7.47	5.66	10.86	10.65	3.07	4.16	4.96	7.07	8.52			-	-
55	NT2RP4001206	64.73	49.7	69.67	51.07	49.14	54.08	37.12	41.91	43.36	*	*	-	-
	NT2RP4001207	7.32	3.58	10.38	7.77	2.4	3.4	3.22	3.79	1.65			-	-
	NT2RP4001210	10.03	7.34	12.93	9.04	5.66	7.56	3.96	6.73	5.83			-	-

Table 454

	NT2RP4000568	23.22	25.79	25.36	10.18	10.21	8.47	12.24	9.65	11.67	**	**	-	-
	NT2RP4000583	5.85	2.98	7.97	5.46	2.06	2.6	3.62	4.57	3				
5	NT2RP4000585	9.33	19	28.37	6.6	3.13	5.64	10.5	7.96	7.04				
	NT2RP4000588	7.43	10.21	10.68	6.68	2.44	1.85	5.44	2.87	5.6	*	*	-	-
	NT2RP4000590	6.55	7.08	8.76	6.12	1.79	2.53	5.82	2.45	2.62		*	-	-
	NT2RP4000599	40.85	42.1	36.12	25.51	13.35	23.08	24.88	16.83	13.91	*	**	-	-
	NT2RP4000603	16.26	11.07	12.79	9.86	5.09	7.92	5.61	11.76	8.02	*		-	-
10	NT2RP4000607	18.04	30.52	24.32	19.53	17.65	25.01	23.84	28.13	28.14				
	NT2RP4000614	10.28	8.25	15.91	8.83	8.89	11.93	10.16	9.95	16.38				
	NT2RP4000634	6.36	4.3	5.27	7.5	2.83	3.42	3.64	2.83	2.88		*	-	-
	NT2RP4000638	23.34	31.48	44.47	17.42	11.63	10.92	16.58	9.75	15.96	*	*	-	-
	NT2RP4000648	20.24	25.67	26.97	19.17	9.45	12.88	31.67	25.66	31.95	*		-	-
	NT2RP4000657	29.17	33.57	37.98	49.46	41.57	42.06	36.04	23.15	32.67	*		+	-
	NT2RP4000691	5.59	4.16	3.63	5.96	2.37	3.18	5.01	2.34	4.68				
15	NT2RP4000697	6.38	8.16	10.53	9.34	4.77	12.75	6.26	6.22	8.11				
	NT2RP4000704	1.95	1.59	2.11	5.91	1.47	3.12	1.95	2.78	2.48				
	NT2RP4000710	20.97	14	15.55	13.68	8.51	18.99	7.53	9.31	11.22	*		-	-
	NT2RP4000713	4.84	3.26	2.49	5.57	2.13	4.31	3.32	1.62	2.08				
	NT2RP4000724	3.62	3.53	2.74	6.31	1.65	2.66	3.33	4.03	5.81				
20	NT2RP4000725	3.89	2.42	3.21	6.35	1.7	3.24	4.75	1.9	3.76				
	NT2RP4000728	11.56	10.9	20.97	12.34	10.22	7.45	16.8	22.52	15.47				
	NT2RP4000737	13.76	11.71	16.21	12.21	6.25	7.06	13.63	12.39	10.01				
	NT2RP4000739	8.74	4.93	10.91	9.95	3.76	6.63	5.46	5.27	7.1				
	NT2RP4000749	16.46	11.21	22.1	13.84	13.31	13.35	10.75	11.02	11.19				
	NT2RP4000769	9.99	10.26	18.79	12.18	3.6	7.13	3.73	5.86	5.05				
25	NT2RP4000774	9.26	8.05	14.72	7.64	2.48	6.01	4.16	4.89	5.59				
	NT2RP4000781	4.81	3.97	6.84	6.54	2.43	3.37	3.39	2.48	3.03				
	NT2RP4000783	17.33	30.73	35.46	25.7	13.51	13.28	15.68	13.86	21.97				
	NT2RP4000788	24.76	10.56	23.27	26.79	17.17	17.48	11.21	13.65	10.18				
	NT2RP4000792	5.04	4.94	10.96	8.67	1.86	4.24	2.65	4.16	5.52				
	NT2RP4000809	342.08	122.2	157.03	126.01	165.45	173.36	118.64	164.8	122.66				
30	NT2RP4000817	8.94	6.91	14.48	7.71	3.01	5.35	5.37	3.63	4.85				
	NT2RP4000821	136.37	90.35	131.55	91.69	114.89	72.17	100.95	73.86	85.69				
	NT2RP4000822	14.55	12.26	20.9	11.12	8.44	15.29	16.17	13.27	18.76				
	NT2RP4000823	2939.8	2567.2	3305.2	1530.8	2690.7	2293.4	1706.4	1550.9	2562.9				
	NT2RP4000831	78.61	54.25	90.69	72.05	94.12	99.31	70.64	58.37	82.02				
	NT2RP4000833	27.77	14.49	23.94	22.19	7.07	21.11	19.12	24.88	29.44				
35	NT2RP4000837	6.72	4.87	9.85	8.02	2.92	3.66	2.57	4.16	7.79				
	NT2RP4000839	145.23	125.82	232.83	173.61	169.29	177.66	124.91	117.96	102.54				
	NT2RP4000846	29.27	18.32	20.88	21.05	17.61	12.28	9.05	14.78	13	*		-	-
	NT2RP4000848	18.2	11.65	15.75	19.15	9.44	13.53	19	14.72	12.39				
	NT2RP4000855	6.13	7.14	11.29	10.66	4.62	6.62	4.65	3.21	3.53				
	NT2RP4000863	6.72	6.6	8.48	8.17	4.81	5.97	4.42	5.29	3.77	**		-	-
40	NT2RP4000865	45.32	26.52	38.54	39.71	38.1	49.51	43.48	35.35	41.09				
	NT2RP4000873	144.06	68.9	135.34	139.48	79.72	155.93	87.22	97.84	116.06				
	NT2RP4000874	10.17	6.06	16.44	7.29	2.79	4.52	3.81	5.9	8.53				
	NT2RP4000875	14.13	12.82	21.34	9.26	7.3	7.14	8.99	11.91	13.22	*		-	-
	NT2RP4000878	36.06	33.12	27.77	33.84	31.46	17.83	20.58	24.01	24.14	*		-	-
	NT2RP4000879	8.38	6.53	9.14	6.87	3.51	3.32	2.85	3.36	3.88	**		-	-
45	NT2RP4000880	35.09	26.71	47.85	50.74	42.76	41.2	29.03	30.06	33.07				
	NT2RP4000891	701.32	715.69	430.85	696.54	840.24	746.75	898.32	324.76	792.55				
	NT2RP4000894	21.41	14.41	21.14	15.13	8.85	12.55	11.08	19.34	17.24				
	NT2RP4000898	2.97	2.72	11.52	4.95	1.01	2.82	2.18	3.29	4.66				
	NT2RP4000899	44.53	41.78	65.07	39.31	37.51	42.09	31.81	35.26	42.89				
	NT2RP4000907	4.19	4.02	10.25	6.24	3.94	2.33	1.69	3.11	0.94				
50	NT2RP4000908	14.45	17.63	18.1	10.04	7.21	5.31	4.66	6.8	4.87	**	**	-	-
	NT2RP4000910	216.22	171.01	317.29	253.37	286.44	240.15	188.21	180.84	183.37				
	NT2RP4000918	3.56	6.94	4.06	7.07	6.01	6.69	5.32	2.08	2.17				
	NT2RP4000925	5.57	5.14	8.28	6.79	4.75	4.89	3.2	5.18	4.02				
	NT2RP4000927	3.1	2.26	4.02	4.89	1.52	0.68	2.5	1.38	1.2				
	NT2RP4000928	9.4	7.34	14.03	9.56	4.86	8.73	4.09	5.17	5.52				
55	NT2RP4000929	3.84	3.91	13.46	5.95	2.85	3.04	2.12	4.29	1.99				
	NT2RP4000946	6.59	6.92	14.08	8.15	4.03	3.77	3.48	3.97	1.29				
	NT2RP4000947	3.36	4.07	10.59	4.84	0.29	0.97	0.36	0.58	0				

Table 453

	NT2RP4000323	42.99	31.25	39.68	23.73	23.52	24.08	25.22	26.31	20.02	*	*	-	-
	NT2RP4000324	28.87	20.72	25.53	23.74	16.73	21.66	28.32	24.57	21.68				
5	NT2RP4000334	11.51	10.77	15.15	9.55	9.47	9.13	6.39	6.12	5.89	**		-	
	NT2RP4000343	20.88	12.62	32.18	18.49	14.22	8.72	14.98	17.89	14.7				
	NT2RP4000348	12.35	14.21	17.96	11.42	8.77	9.63	12.1	7.85	8.02				
	NT2RP4000349	8.77	16.24	12.89	11.44	12.7	9.21	11.37	7.34	11.23				
	NT2RP4000355	253.25	117.6	189.34	264.37	149.78	254.47	257.02	316.08	192.79				
	NT2RP4000356	61.98	36.7	79.91	60.03	57.95	82.79	39.95	40.28	37.12				
10	NT2RP4000360	38.52	30.1	36.81	28.77	30.27	34.86	22.5	29.04	23	*		-	
	NT2RP4000367	5.32	7.72	13.11	9.15	7.75	6.58	5.65	7.52	4.05				
	NT2RP4000370	6.13	3.8	7.29	7.4	4.81	4.6	4.29	3.27	2.44				
	NT2RP4000373	52.16	35.56	49.83	33.3	22.86	20.18	8.95	13.1	7.55	*	**	-	-
	NT2RP4000376	4.35	4.76	3.85	5.98	2.55	3.6	3.64	2.24	1.76	*		-	
	NT2RP4000381	4.21	7.62	7.46	8.53	4.41	5.48	6.16	4.44	4.11				
15	NT2RP4000388	15.3	8.64	28.13	9.56	6.37	8.78	3.98	7.28	10.59				
	NT2RP4000390	9.32	12.91	21.1	9.57	7.3	6.9	4.45	8	7.73				
	NT2RP4000393	4.73	6.04	14.2	5.91	3.76	3.4	2.49	4.28	1.61				
	NT2RP4000398	15.1	12.32	17.4	15.43	7.99	12.09	10.09	7.98	7.95	*		-	
	NT2RP4000406	6.93	7.1	10	7.22	4.81	7.48	4.73	7.43	4.16				
	NT2RP4000407	14.2	8.65	18.88	9.39	7.38	7.23	8.05	7.66	8.08				
20	NT2RP4000413	6.36	8.44	8.74	7.32	6.23	4.25	14.9	11.64	8.91				
	NT2RP4000415	5.58	6.36	5.25	8.19	3.66	6.69	6.61	5.43	4.11				
	NT2RP4000417	4.81	3.78	14.82	6.27	2.89	2.96	1.78	3.56	2.97				
	NT2RP4000423	8.56	11.85	52.11	7.59	6.28	6.14	5.59	8.25	4.46				
	NT2RP4000424	5.63	6.45	12.56	7.04	3.08	2.9	3.66	4.14	1.14				
25	NT2RP4000447	7.22	11.02	14.44	6.8	2.54	3.42	2.02	2.94	1.68	*		-	
	NT2RP4000448	10.44	4.78	12.49	7.44	5.19	4.9	2.34	3.11	4.18				
	NT2RP4000449	14.23	13.95	15.79	8.35	8.81	8.15	9.09	11.62	7.03	**	*	-	-
	NT2RP4000453	9.01	6.07	14.02	7.74	5.15	4.89	6.56	4.27	8.08				
	NT2RP4000455	9.03	10.34	8.68	7.99	5.45	3.17	4.44	4.36	4.13	**		-	
	NT2RP4000456	7.04	6.17	17.02	5.76	3.52	4.72	1.66	2.58	3.46				
30	NT2RP4000457	13.26	8.71	17.6	8.13	4.23	7.52	11.15	9.14	8.56				
	NT2RP4000461	55.44	37.43	46.46	34.13	24.48	24.96	29.53	26.44	29.5	*	*	-	-
	NT2RP4000462	42.85	30.75	56.94	75.69	50.54	40.64	52.98	43.57	42.53				
	NT2RP4000463	85.58	62.21	92.77	65.01	52.89	43.38	67.48	75.6	83.4				
	NT2RP4000471	25.28	34.12	24.81	15.24	25.16	15.87	19.6	16.58	11.2	*		-	
	NT2RP4000472	7.71	9.85	9.52	10.37	5.54	5.62	6.84	6.67	3.82				
35	NT2RP4000476	14.81	18.77	16.55	11.79	4.98	3.82	5.79	4.35	2.67	*	**	-	-
	NT2RP4000480	12.42	10.39	22.37	9.71	8.11	7.69	2.88	6.14	7.56				
	NT2RP4000481	7.17	5.4	16.93	6.71	1.97	4.16	2.54	3.59	3.12				
	NT2RP4000483	21.87	12.5	29.34	14.63	7.41	9.82	17.08	21.67	14.93				
	NT2RP4000487	11.87	8.84	15.81	9.26	6.43	6.73	8.06	10.31	7.08				
	NT2RP4000488	16.33	41.67	19.05	11.91	7.25	5.15	7.41	10.6	17.63				
40	NT2RP4000487	5.81	3.81	10.08	3.87	3.09	2.81	2.48	1.77	2.22				
	NT2RP4000488	26.54	11.6	46.3	21.94	11.56	20.83	11.71	6.48	8.74				
	NT2RP4000500	16.19	14.2	13.24	21.4	11.25	11.2	15.02	6.9	5.02				
	NT2RP4000507	14.47	9.74	21.49	11.59	12.91	17.98	12.44	18.01	18.95				
	NT2RP4000515	34.88	29.88	36.21	23.73	25.4	27.6	24.2	28.54	29.19	*		-	
	NT2RP4000516	8.8	4.78	15.85	7.22	3.73	6.26	5.56	4.23	5.23				
45	NT2RP4000517	6.93	4.39	11.92	5.6	2.28	2.95	2.26	2.51	3.47				
	NT2RP4000518	5.66	5.23	9.62	6.13	3.29	2.14	2.66	2.57	2.25	*		-	
	NT2RP4000519	11.25	13.8	9.43	7.22	5.12	6.11	9.96	4.77	2.74	*		-	
	NT2RP4000524	3.66	2.61	3.86	4.66	0.22	1.66	1.44	0.06	0.23	**		-	-
	NT2RP4000528	9.85	11.61	11.68	7.98	5.49	4.94	7.65	1.72	3.24	*	*	-	-
	NT2RP4000537	22.62	26.04	23.49	9.39	8.19	12.07	6.5	9.43	12.67	**	**	-	-
50	NT2RP4000541	6.77	5.01	14.31	6.57	5.43	3.77	2.53	4.6	6.37				
	NT2RP4000543	7.3	6.98	11.29	8.27	7.56	3.29	6.55	3.66	5.22				
	NT2RP4000545	22.52	30.13	32.57	13.3	16.56	19.62	18.71	21.42	13.96	*	*	-	-
	NT2RP4000546	4.78	4.57	9.21	6.74	1.72	3.53	3.64	0.82	4.41				
	NT2RP4000549	11.61	18.79	20.88	6.99	2.21	3.33	4.79	2.09	3.96	*	*	-	-
	NT2RP4000556	15.54	16.2	17.27	7.73	3.7	5.61	5.6	2.76	4.84	**	**	-	-
55	NT2RP4000557	11.07	12.42	10.89	8.61	4	8.57	12.02	7.59	12.85				
	NT2RP4000558	14.08	10.84	18.04	10.75	6.08	12.87	11.5	16.9	20.52				
	NT2RP4000560	6.31	4.01	7.32	5.77	2.76	5.36	5.37	6.37	4.62				

Table 452

	NT2RP3004670	5.78	5.9	15.55	5.95	8.77	4.78	1.73	4.19	3.14			
	NT2RP4000008	4.28	3.76	9.74	7.01	3.13	2.8	0.87	1.61	1.16			
5	NT2RP4000018	777.26	720.79	779.71	458.85	552.02	180.81	608.14	341.39	573.73	*	*	-
	NT2RP4000023	12.08	13.49	11.95	6.59	4.27	3.03	5.43	2.46	2.58	**	**	-
	NT2RP4000025	10.92	10.01	10.19	8.87	3.46	6.52	5.55	3.38	7.2	*	*	-
	NT2RP4000035	13.64	12.06	18.1	13.28	10.2	10.62	12.01	14.28	15.16			
	NT2RP4000041	7.95	4.85	12.17	6.47	2.4	5.6	2.83	2.55	3.55			
	NT2RP4000049	5.5	8.69	14.31	6.89	4.49	2.04	3	3.25	4.73			
10	NT2RP4000050	7.92	6.2	13.4	9.91	4.4	3.61	2.52	5.2	2.39			
	NT2RP4000051	11.64	12.9	17.08	9.37	6.75	7.7	7.83	11.79	8.18	*	*	-
	NT2RP4000063	9.36	7.43	9.4	6.21	2.66	4.7	2.12	2.88	2.29	*	**	-
	NT2RP4000065	8.57	5.45	7.62	5.43	1.38	3.74	3.79	2.44	1.14	*	*	-
	NT2RP4000070	8.9	10.6	11.32	10.92	5.17	4.74	3.9	3	5.41	**	**	-
	NT2RP4000074	7.09	6.56	7.33	6.07	3.64	2.14	2.7	1.95	3.11	**	*	-
15	NT2RP4000078	10.64	7.98	15.22	9.04	4.02	9.72	4.12	7.85	6.44			
	NT2RP4000080	39.53	29.48	35.07	14.73	7.78	7.51	7.13	12.84	8.56	**	**	-
	NT2RP4000099	17.27	15.29	21.59	9.54	12.81	6.62	5.87	6.72	3.64	*	**	-
	NT2RP4000102	8.52	8.93	15.05	7.45	6.98	5.31	5.6	5.59	5.12			
	NT2RP4000103	20.66	29.01	31.17	15.7	16.08	13.47	20.04	13.58	12.68	*	*	-
	NT2RP4000108	10.35	7.66	11.54	5.11	5.39	5.22	4.96	2.69	3.44	*	**	-
20	NT2RP4000109	9.73	11.84	8.78	8.41	4.9	6.21	6.7	3.67	4.98	*	*	-
	NT2RP4000111	9.68	7.22	7.83	10.68	5.18	7.24	9.51	4.5	2.88			
	NT2RP4000112	16.91	11.4	24.28	20.79	19.36	22.3	16.09	13.49	17.44			
	NT2RP4000115	7.34	4.25	10.98	5.53	2.81	4.32	2.76	2.51	2.27			
	NT2RP4000129	24.84	25.48	33.51	13.19	12.49	13.21	12.38	15.5	12.9	**	**	-
	NT2RP4000137	20.7	16.54	14.08	10.11	6.13	6.89	9.81	6.38	10.47	*	*	-
25	NT2RP4000138	7.91	6.25	8.72	5.36	2.95	3.73	3.09	3.11	5.54	*	*	-
	NT2RP4000141	11.63	7.1	7.39	5.05	2.2	2.55	3.83	2.34	1.67	*	*	-
	NT2RP4000147	7.8	8.75	6.46	5.67	2.45	1.71	3.71	1.66	2.95	*	**	-
	NT2RP4000150	22.83	40.43	32.64	26.18	18.45	19.86	24.1	13.63	20.82			
	NT2RP4000151	5.35	2.77	7.92	5.7	2.57	4.58	2.81	1.73	3.64			
	NT2RP4000157	5.03	3.17	7.55	5.12	2.41	3.03	2.61	4.59	2.88			
30	NT2RP4000159	7.7	6.21	14	7.81	5.43	9.16	5.02	7.93	7.16			
	NT2RP4000163	11.36	16.35	22.24	7.82	6.7	5.1	11.58	8.66	14.17	*	*	-
	NT2RP4000167	5.72	10.76	11.28	6.86	1.93	2.94	6.09	1.64	2.98			
	NT2RP4000171	17.67	20.8	16.76	13.24	7.81	11.48	17.22	11.6	13.85	*	*	-
	NT2RP4000175	7.68	11.5	9.21	7.19	5.94	7.44	5.84	4.86	6.71	*	*	-
35	NT2RP4000180	8.25	9.87	9.45	8.5	4.86	7.21	9.47	6.55	7.72			
	NT2RP4000185	3.44	9.69	9.07	5.25	3.82	5.08	3.89	5.51	9.51			
	NT2RP4000192	3.66	2.3	7.85	7.16	4.58	3.22	2.99	2.89	2.84			
	NT2RP4000194	4.94	3.92	6.27	5.67	6.92	4.33	4.73	3.32	6.3			
	NT2RP4000196	4.44	3.86	8.22	4.42	2.19	1.89	2.14	2.44	1.01			
	NT2RP4000210	4.32	3.9	5.46	5.03	2.17	1.97	3.56	2.52	3.91			
40	NT2RP4000212	10.41	10.12	9.74	6.43	6.54	3.07	4.86	4.69	2.44	*	**	-
	NT2RP4000214	21.41	28.56	31.47	14.93	12.43	12.75	14.71	10.53	11	*	**	-
	NT2RP4000216	18.56	17.54	17.8	9.05	3.32	6.3	7.12	4.66	4.69	**	**	-
	NT2RP4000218	20.76	13.22	21.73	16.85	12.17	21.91	12.16	18.58	23.33			
	NT2RP4000223	11.85	6.5	8.31	9.5	4.18	9.75	8.04	7.05	11.84			
	NT2RP4000243	10.56	9.93	12.96	11.2	9.86	11.7	8.64	9.69	10.23			
45	NT2RP4000246	7.12	5.86	5.64	5.65	4.43	4.42	2.37	2.31	3.27	**	*	-
	NT2RP4000250	11.29	8.98	12.39	10.04	3.49	5.45	5.46	5.81	5.67	**	*	-
	NT2RP4000256	3.26	5.13	6.67	4.51	2.18	2.56	3.11	2.41	2.48			
	NT2RP4000257	23.54	23.13	30.8	34.24	27.3	37.51	26.44	19.58	27.22			
	NT2RP4000259	5.19	5.55	3.64	8.67	2.13	4.17	4.34	3.84	5.01			
	NT2RP4000261	4.72	2.05	7.37	6.8	1.48	3.76	2.37	3.23	3.88			
50	NT2RP4000262	2.93	2.46	11.12	5.21	1.63	2.61	2.49	2.62	2.71			
	NT2RP4000263	8.03	5.82	13.8	7.33	4.75	4.76	8.35	4.63	5.93			
	NT2RP4000280	36.69	25.69	34.43	21.81	13.93	11.89	23.3	25.5	21.75	*	*	-
	NT2RP4000286	7.96	5.71	10.34	8.48	4.4	5.47	4.56	4.72	2.93			
	NT2RP4000290	6.28	5.27	10.35	7.67	3.77	4.58	5.45	2.24	4.35			
	NT2RP4000291	23.37	31.04	45.88	29.65	20.44	28.9	16.58	13.08	6.43	*	*	-
55	NT2RP4000301	9.06	21.31	16.13	18.84	21.42	12.52	13.04	8.64	8.86			
	NT2RP4000312	7.37	5.86	10.71	6.75	4.45	5.45	4.08	4.23	5.32			
	NT2RP4000321	14.11	6.64	13.51	10.77	5.65	7.11	6.38	8.75	12.76			

Table 451

	NT2RP3004338	20.15	16.11	15.61	11.35	11.45	16.93	14.61	18.8	16.46			
	NT2RP3004341	45.52	46.06	66.9	30.5	32.86	24.48	35.75	44.1	42.23	*	-	
5	NT2RP3004345	5.41	7.27	12.03	7.28	2.26	2.19	5.01	3.58	4.59			
	NT2RP3004348	8.2	11.9	19.15	6.37	5.32	6.29	9.92	4.97	6.08			
	NT2RP3004349	46.27	44.89	52.77	25.21	14.4	28.82	51.32	39.58	50.42	**	-	
	NT2RP3004355	13.69	17.95	13.45	12.62	8.18	11.72	10.41	5.58	7.23	*	-	
	NT2RP3004356	6.56	4.92	9.79	7.46	3.18	5.93	2.87	5.09	3.61			
	NT2RP3004360	11.41	7.43	12.2	11.26	4.38	7.34	4.62	4.39	7.23	*	-	
10	NT2RP3004361	22.43	16.96	27.29	12.91	14.7	16.73	18.55	21.02	26.27			
	NT2RP3004374	6.08	4.11	5.92	5.52	2.91	4.87	2.31	2.56	2.6	*	-	
	NT2RP3004378	50.03	39.45	48.92	39.78	39.18	55.24	61.97	46.46	63.83			
	NT2RP3004399	25.24	30.84	27.39	23.98	13.72	17.1	23.05	19.32	24.46	*	-	
	NT2RP3004405	5.46	2.89	3.95	6.86	3.8	2.54	2.58	3.57	3.49			
	NT2RP3004406	48.02	80.1	76.06	27.26	18.49	29.22	40.17	40.49	25.13	*	*	-
15	NT2RP3004411	21.84	13.45	15.39	11.3	8.55	12.78	6.83	10.8	17.8			
	NT2RP3004424	12.83	9.11	14	10.53	6.29	12.31	8.48	6.86	11.07			
	NT2RP3004428	9.06	4.59	7.61	7.92	3	6.19	2.13	2.45	2.73	*	-	
	NT2RP3004432	41.77	40.01	43.16	46.87	29.7	37.75	43.11	55.91	45.48			
	NT2RP3004434	6.59	6.36	9.2	19.42	10.09	11.41	8.35	8.97	10.87			
	NT2RP3004446	5.55	1.31	2.23	5.8	1.27	2.52	2.84	1.81	2.66			
20	NT2RP3004451	28.05	27.38	35.9	43.8	34.57	29.99	27.84	20.66	27.06			
	NT2RP3004454	3.04	4.95	3.07	5.17	2.65	2.97	3.32	1.95	4.76			
	NT2RP3004466	17.88	12.48	17.63	17.19	13.76	19.32	8.38	10.38	11.73	*	-	
	NT2RP3004470	6.58	4.48	12.1	9.37	2.81	3.83	6.66	6.76	5.57			
	NT2RP3004472	4.96	8.32	13.71	7.25	5.54	4.58	2.83	3.65	3.08			
	NT2RP3004475	9.27	6	11.18	8.66	2.73	3.48	3.79	5.2	3.34	*	-	
25	NT2RP3004480	22.33	15.69	32.46	17.18	10.83	15.46	12.05	10.67	10.95			
	NT2RP3004481	7.51	6.36	10.91	7.96	3.04	6.85	8.17	5.21	5.83			
	NT2RP3004490	1.96	3.67	4.31	5.07	2.67	3.48	2.86	2.05	1.78			
	NT2RP3004496	29.05	43.68	52.3	32.37	24.7	28.45	17.59	12.79	20.95	*	-	
	NT2RP3004498	6.1	6.29	10.2	8.29	5.15	5.43	3.36	4.35	5.77			
	NT2RP3004503	13.34	8.17	15.98	13.28	6.98	13.95	9.69	13.04	16.21			
30	NT2RP3004504	26.86	13.87	24.17	16.13	11.34	13.03	8.55	9.71	9.3	*	-	
	NT2RP3004505	60.88	42.22	65.16	60.93	61.18	59.93	38.93	40.66	35.92			
	NT2RP3004507	6.96	5.95	8.15	5.81	3.02	3.27	2.74	2.78	2.24	**	-	
	NT2RP3004519	3.09	3.32	8.34	4.29	1.92	0.77	1.76	0.64	1.58			
	NT2RP3004524	22.27	16.91	24.46	12.8	14.38	10.78	10.59	7.77	17.02	*	-	
	NT2RP3004527	31.26	36.6	41.52	32.19	22.11	22.1	23.08	21.05	18.57	**	-	
35	NT2RP3004534	1954.5	1225.4	1438.4	1526.7	1505.7	1779.6	777.34	2507.5	2418.6			
	NT2RP3004539	4.71	4.69	10.04	5.61	3.18	4.71	4.15	5.52	6.25			
	NT2RP3004541	14.72	10.14	22.05	12.49	7.26	9.05	8.78	11	10.96			
	NT2RP3004546	17.32	13.37	16.21	16.43	8.76	13.79	6.55	10.61	10.66	*	-	
	NT2RP3004551	12.43	7.15	13.73	9.94	5.53	6.65	10.18	9.42	7.62			
	NT2RP3004552	19.37	11.98	21.3	18.2	12.57	16.98	18.98	16.22	18.63			
40	NT2RP3004557	11.18	12.99	15.15	12.15	8.14	12.54	12.08	12.56	14.07			
	NT2RP3004561	2.52	4.37	4.66	6.9	2.44	3.94	3.97	3.31	3.2			
	NT2RP3004566	16.46	13.14	29	10.83	7.99	8.15	3.35	6.43	21.27			
	NT2RP3004569	20.41	16.28	28.7	14.13	11.71	13.43	13.97	16.78	16.75			
	NT2RP3004572	12.48	13.38	19.74	11.19	9.39	9.67	3.98	9.35	8.35	*	-	
	NT2RP3004578	6.49	6.06	8.06	7.21	4.59	4.3	5.13	4.76	4.1	*	-	
45	NT2RP3004584	8.34	6.1	9.57	5.07	2.29	4.9	3.85	3.5	2.08	*	*	-
	NT2RP3004588	4.64	4.62	7.46	7.78	2.88	3.22	3.01	5.54	3.28			
	NT2RP3004594	18.77	25.61	21.44	21.59	19.15	22.94	29.86	27.19	18.84			
	NT2RP3004603	4.79	6.26	5.77	8.8	3.9	4.9	5.46	7.72	5.82			
	NT2RP3004612	6.46	6.92	12.58	12.71	4.69	5.32	2.31	10.42	14.75			
50	NT2RP3004617	8.05	8.14	15.39	8.87	3.8	5.79	3.78	6.88	7.16			
	NT2RP3004618	36.56	23.11	37.29	24.87	25.7	22.47	22.39	33.72	29.91			
	NT2RP3004625	19.69	16.42	18.21	11.76	10.91	14.71	6.25	8.76	11.12	*	**	-
	NT2RP3004635	11.6	9.72	13.73	7.06	5.4	4.21	3.68	2.39	2.94	*	**	-
	NT2RP3004640	4.25	10.31	12.87	7.49	7.74	5.34	4.28	3.2	9.33			
	NT2RP3004642	15.02	23.4	27.93	15.96	15.92	16.39	17.49	17.88	14.1			
55	NT2RP3004647	10.52	11.7	14.89	13.69	6.27	6.34	18.22	14.86	18.18	*	*	+
	NT2RP3004652	28.1	24.38	41.76	13.95	14.47	12.02	4.02	12.27	12.65	*	*	-
	NT2RP3004669	7.74	7.41	20.24	7.31	6.69	6.49	4.02	6.68	4.21			

Table 450

	NT2RP3004005	23.75	15.81	26.46	14	10.64	14.12	16.28	18.58	18.43			
	NT2RP3004013	22.54	11.33	21.98	6.11	3.13	3.73	3.05	4.94	3.43	*	*	-
5	NT2RP3004016	7.39	7.18	13.3	10.92	9.82	5.35	6.27	5.44	4.76			
	NT2RP3004025	22.82	15.93	25.25	22.37	10.03	12.12	21.2	17.91	21.7			
	NT2RP3004030	4.66	4.65	8.61	4.83	3.44	3.8	4.36	2.41	1.96			
	NT2RP3004041	4.1	2.26	3.06	5.03	2.28	1.57	2.62	0.7	0.71			
	NT2RP3004042	2602.1	1724.7	1778.2	2098.3	1690	1795.1	1947.3	2828.9	2085.3			
	NT2RP3004044	32.34	21.11	30.01	12.91	11.29	11.35	7.18	15.29	26.2	**		-
10	NT2RP3004051	19.22	22.27	34.06	33.29	21.4	25.83	12.31	39.18	27.89			
	NT2RP3004052	20.9	17.89	21.89	18.39	15.57	16.97	13.27	17.39	19.16			
	NT2RP3004053	19.14	11.65	20.87	15.13	10.18	12.84	3.1	7.99	5.77	*		-
	NT2RP3004055	5.06	6.81	8.6	5.35	3.54	2.7	2.8	3.03	2.13	*		-
	NT2RP3004059	26.03	28.01	36.98	19.3	19.54	21.1	20.16	18	15.03	*	*	-
	NT2RP3004063	29.14	22.42	34.14	23.83	23.28	19.51	31.01	26.64	12.72			
15	NT2RP3004067	7.95	8.62	20.76	10	4.39	4.65	3.75	4.12	8.39			
	NT2RP3004070	905.61	934.59	965.28	291.19	622.85	595.78	278.31	1360.6	1279.5	*		-
	NT2RP3004075	11.59	15.27	23.01	8.7	5.82	8.89	3.95	5.77	5.73	*		-
	NT2RP3004078	13.83	11.53	17.13	13.53	5.08	9.69	12.68	11.48	12.45			
	NT2RP3004083	33.39	27.25	43.84	12.1	13.49	7.53	6.38	6.07	6.79	*	**	-
	NT2RP3004084	25.34	24.44	39.2	27	15.25	16.2	38.67	47.1	47.33			
20	NT2RP3004087	13.47	17.06	17.12	15.44	8.85	8.53	10.22	10.37	8.13	*		-
	NT2RP3004090	72.24	81.09	78.4	57.09	57.97	55.36	77.78	70.88	90.98	**		-
	NT2RP3004093	28.49	18.68	45.95	14.41	12.09	18.33	8.92	23.19	25.5			
	NT2RP3004095	33.26	26.66	48.11	23.16	21.66	33.49	19.24	23.44	21.47			
	NT2RP3004102	34.73	35.32	40.77	25.98	17.76	24.22	15.74	23.36	12.18	*	**	-
	NT2RP3004110	7.05	4.57	10.07	7.67	1.49	3.89	2.25	3.65	1.62	*		-
25	NT2RP3004119	11.05	7.96	13.76	6.77	3.42	4.11	3.84	3.73	1.51	*	*	-
	NT2RP3004125	3.06	3.32	9.52	6.05	1.93	1.78	1.81	1.49	1			
	NT2RP3004129	11.05	13	16.54	13.66	11.08	8.18	7.67	3.97	5.87	*		-
	NT2RP3004130	6.85	5.7	8.22	6.11	4.3	3.04	1.91	3.1	2.41	**		-
	NT2RP3004133	8.31	5.24	20.22	8.3	3.21	4.09	3.33	4.08	3.29			
	NT2RP3004145	5.99	4.29	13.16	5.33	2.78	2.69	2.63	4.11	3.14			
30	NT2RP3004148	53	37.5	48.88	32.7	23.92	15.17	30.82	36.4	44.13	*		-
	NT2RP3004155	8.18	8.86	12.7	12.14	3.88	4.25	2.59	2.84	3.13	**		-
	NT2RP3004165	77.97	53.41	68.62	51.37	46.12	31.89	49.51	25.55	55.19			
	NT2RP3004179	9.55	6.18	11.2	6.95	2.94	2.88	4.01	3	3.7	*		-
	NT2RP3004185	6.31	5.21	6.12	9.04	2.95	2.38	1.55	0.61	2.3	**		-
	NT2RP3004188	6.31	7.01	6.86	7.94	4.08	2.67	3.74	2.3	1.82	**		-
35	NT2RP3004189	15.63	7.3	21.16	9.58	5.21	8.89	4.83	8.4	6.14			
	NT2RP3004190	64.52	34.39	57.67	48.45	40.38	53.55	47.37	42.21	45.49			
	NT2RP3004191	122.58	123.5	136.79	74.78	46.86	44.19	48.51	70.16	47.4	**	**	-
	NT2RP3004202	18.55	13.31	21.41	14.01	15.47	6.42	10.97	12.13	10.67			
	NT2RP3004205	10.59	7.33	14.79	6.1	4.73	2.92	2.52	2.35	2.6	*		-
40	NT2RP3004206	15.55	12.16	17.94	7.47	8.64	7.63	5.69	7.13	12.18	*		-
	NT2RP3004207	8.65	8.42	9.24	5.53	2.88	3.67	2.31	1.22	2.02	**	**	-
	NT2RP3004209	27.4	23.65	14.82	17	13.67	16.8	19.95	17.39	12.49			
	NT2RP3004215	22.45	23.53	27.86	11	7.75	10.82	4.22	8.07	7.54	**	**	-
	NT2RP3004219	5.58	5.12	13.51	6.61	2.08	4.81	3.19	3.17	3.09			
	NT2RP3004242	4.18	6.26	13.02	6.15	3.05	4.41	4.72	3.98	3.45			
	NT2RP3004246	3.85	4.1	9.37	4.89	2.24	1.94	1.48	3.04	2.11			
45	NT2RP3004253	7.46	3.55	8.07	5.38	2.69	1.44	2.57	2.47	3.21			
	NT2RP3004258	8.43	4.33	9.6	7.18	4.84	3.38	6.14	4.76	3.41			
	NT2RP3004262	5.75	6.49	6.12	6.38	2.36	2.74	4.69	0.6	2.65	*		-
	NT2RP3004275	5.37	6.56	6.47	6.33	1.33	2.3	2.5	1.89	3.75	**		-
	NT2RP3004282	4.86	27.86	18.63	6.39	3.74	5.05	2.71	3.54	3.27			
50	NT2RP3004289	6.43	5.13	10.92	5.92	2.79	3.35	2.25	2.59	3.21			
	NT2RP3004294	4.71	4.44	6.56	4.95	2.62	3.32	2.88	6.45	3.88			
	NT2RP3004298	64.72	55.93	73.74	63.66	58.3	70.43	70.1	51.46	68.44			
	NT2RP3004309	10	7.69	11.87	7.21	2.5	3.75	5.84	3.48	5.36	*	*	-
	NT2RP3004321	23.29	38.14	42.57	12.44	16.53	8.76	14.14	5.43	9.71	*	*	-
	NT2RP3004322	4.19	4.29	4.98	4.8	1.3	0.91	2.93	1.84	2.18	**		-
55	NT2RP3004332	9.53	11.83	11.13	7.92	3.85	4.05	4.3	2.24	2.92	*	**	-
	NT2RP3004334	14.9	13.13	17.93	15.28	11.06	7.25	6.62	11.71	9.43	*		-
	NT2RP3004336	9.94	8.5	13.66	9.26	7.37	6.68	4.73	8.17	4.32			

Table 449

	NT2RP3003716	7.01	8.27	14.18	5.47	2.75	3.52	6.58	5.29	3.83			
	NT2RP3003721	13.04	10.15	23.8	10.64	5.81	9.58	11.26	11.19	10.89			
5	NT2RP3003722	3.89	8.42	15.57	8.33	6.34	5.67	3.09	6.47	3.17			
	NT2RP3003726	9.91	10.2	12.37	9.7	5.8	5.18	6.04	11.22	5.51			
	NT2RP3003729	15.24	10.41	15.84	9.27	8.89	4.46	6.77	7.5	6.93	*		-
	NT2RP3003731	21.75	24.76	24.08	12.03	11.65	6.88	4.57	8.08	5.08	**	**	-
	NT2RP3003740	6.25	8.21	13.85	9.2	5.84	6.01	5.86	5.22	4.61			
	NT2RP3003746	8.07	4.76	18.45	7.1	1.15	4.58	3.94	4.56	4.9			
10	NT2RP3003749	12.41	10.83	19.25	6.47	8	6.02	6.17	7.75	8.42			
	NT2RP3003754	6.51	6.89	16.88	6.39	6.73	2.6	13.69	10.54	7.77			
	NT2RP3003759	5.04	4.82	10.4	5.36	1.64	1.95	2.96	2.71	1.17			
	NT2RP3003764	37.79	25.83	53.35	42.39	40.22	42.13	25.68	27.55	28.87			
	NT2RP3003766	53.84	52.07	31.23	39.67	26.34	21.22	14.6	22.88	28.5	*		-
	NT2RP3003767	8.01	8.08	7.88	8.43	4.34	4.24	4.39	2.88	4.34	**		-
15	NT2RP3003778	24.16	27.96	20.47	21.14	16.84	15.25	32.15	22.31	26.7			
	NT2RP3003779	12.68	5.29	15.64	10.68	4.35	10.53	3.1	5.95	4.41			
	NT2RP3003783	21.6	12.18	19.59	7.99	4.9	7.05	6.28	6.7	5.9	*	*	-
	NT2RP3003787	6.7	5.75	13.41	6.04	3.64	1.85	4.2	4.99	1.46			
	NT2RP3003789	8.11	11.36	13.5	7.92	6.55	8.67	5.52	5.69	3.28	*		-
	NT2RP3003795	28.23	17.85	14.78	13.94	17.1	12.18	13.09	11.24	13.99			
20	NT2RP3003799	7.73	3.87	9.64	5.67	2.6	3.17	2.83	1.93	2.55			
	NT2RP3003800	80.77	76.75	98.8	62.18	49.64	73.93	101.61	58.63	58.22			
	NT2RP3003805	19.72	20.08	15.07	17.29	8.51	14.67	17.48	12.39	11.94			
	NT2RP3003809	8.63	14.84	28.37	7.81	4.06	5.86	6.78	7.04	6.7			
	NT2RP3003818	19.91	12.71	21.8	8.29	12.89	15.12	8.35	11.58	14.32			
25	NT2RP3003824	4.44	5.34	16.42	6	3.26	2.18	5.16	4	3.14			
	NT2RP3003825	15.83	8.99	17.57	10.52	9.35	8.99	8.88	7.14	8.67			
	NT2RP3003828	7.18	6.66	13.27	7.4	5.95	4.81	5.75	3.03	3.24			
	NT2RP3003831	49.77	38.05	46.92	12.19	13.59	13	11.7	8.2	8.68	**	**	-
	NT2RP3003833	7	12.76	5.52	6.51	1.52	3.02	3.01	1.46	3.08			
	NT2RP3003836	22.92	45.16	37.16	16	14.78	18.49	26.73	7.91	11.58	*		-
30	NT2RP3003842	3.09	1.45	4.55	4.66	0.91	2.25	1.5	2.49	0.97			
	NT2RP3003843	23.17	13.66	18.94	21.62	11.25	18.44	16.81	22.17	19.21			
	NT2RP3003844	11.64	8.51	12	9.56	4.9	7.1	7.94	10.02	13.51			
	NT2RP3003846	11.56	8.88	8.25	8.42	3.89	7.04	8.15	6.21	10.88			
	NT2RP3003849	5.16	5.61	9.2	5.16	1.5	2.43	2.36	2.09	1.97	*		-
	NT2RP3003862	8.43	6.43	11.03	6.94	2.53	2.38	2.84	2.47	3.72	*		-
35	NT2RP3003870	9.39	14.91	15.6	7.42	3.58	2.61	4.29	2.22	5.4	*	*	-
	NT2RP3003874	8.27	8.06	7.54	6.09	3.19	3.65	2.59	2.64	5.21	*	**	-
	NT2RP3003876	6.84	5.28	7.66	10.08	2.52	6.72	2.11	6.22	4.3			
	NT2RP3003880	11.2	8.02	13.91	8.47	2.77	7.12	3.7	8.83	6.95			
	NT2RP3003889	6.9	11.63	11.35	8.75	7.41	9.51	7.37	8.06	5.22			
	NT2RP3003891	13.51	11.29	17.94	8.42	7.04	7.49	10.43	12.21	12.8	*		-
40	NT2RP3003914	2.72	1.62	6.58	5.33	2.56	2.54	2.7	1.4	2.69			
	NT2RP3003915	13.41	9.97	16.94	9.16	3.98	5.29	7.39	4.87	7.36	*	*	-
	NT2RP3003918	7.35	6.41	5.09	8.06	2.84	2.93	4.26	2.94	4.43	*		-
	NT2RP3003920	10.75	6.92	10.79	7.41	1.85	6.22	6.71	4.36	3.03	*		-
	NT2RP3003924	33.52	17.52	18.55	12.63	9.6	13.46	9.41	16.84	13.76			
	NT2RP3003932	13.16	5.83	7.58	6.21	2.22	4.38	1.9	2.22	2.59	*		-
45	NT2RP3003939	11.85	14.76	15.54	9.97	7.98	8.34	3.34	3.94	3.84	*	**	-
	NT2RP3003940	12.22	9.19	12.29	11.49	8.56	4.42	7.78	12.51	10.79			
	NT2RP3003943	7.2	4.68	6.57	7.12	5.58	5.18	4.42	3.92	6.05			
	NT2RP3003959	7.72	13.55	11.27	9.14	1.71	5.84	6.69	7.88	9.75			
	NT2RP3003963	75.69	75.77	72.03	129.26	82.85	106.1	81.78	51.98	65.78			
	NT2RP3003965	18.74	14.24	13.95	14.48	9.03	12.62	20.67	19.68	26.47			
50	NT2RP3003972	4.57	5.37	14.74	8.17	2.14	5.46	5	3.91	5.17			
	NT2RP3003973	6.48	6.48	12.61	7.2	3.25	3.55	3.26	4.26	5.78			
	NT2RP3003979	4.45	3.72	9.63	4.82	2.41	2.62	1.15	1.72	1.66			
	NT2RP3003980	146.06	117.79	158.74	103.46	88.31	102.32	155.94	132.47	147.6	*		-
	NT2RP3003982	7.33	5.76	16.01	6.18	2.48	3	4.61	2.79	5.02			
	NT2RP3003989	10.71	10.02	27.87	8.13	4.12	8.61	5.84	9.52	13.56			
55	NT2RP3003992	46.04	43.58	55.68	31.62	22.83	36.4	32.87	30.71	18.85	*	*	-
	NT2RP3004000	75.38	38.37	57.12	93.25	54.67	51.6	36.78	36.39	33.4			
	NT2RP3004001	14.94	10.16	22.03	15.95	8.73	15.13	13.16	15.23	21.06			

Table 448

	NT2RP3003396	16.4	12.74	16.94	16.32	6.79	16.93	14.73	19.34	18.49			
	NT2RP3003403	3.1	2.78	7.09	5.18	1.78	3.29	1.96	3.43	3.41			
5	NT2RP3003409	7.35	11.81	12.63	9.44	12.96	4.77	6.95	7.18	6.37			
	NT2RP3003411	11.06	12.25	14.86	7.42	4.83	5.8	7.12	7.74	6.44	**	**	-
	NT2RP3003420	19.06	32.9	36.55	21.71	11.78	14.57	23.43	20.87	21.3			
	NT2RP3003425	6.27	5.68	8.42	5.84	2.62	3.03	4.29	2.39	4.38	*		-
	NT2RP3003426	7.48	8.53	6.3	7.22	3.59	4.34	6.93	6.05	8.2			
10	NT2RP3003427	28.71	32.12	28.4	30.38	20.54	27.6	31.57	21.98	21.07			
	NT2RP3003433	4.33	3.61	3.8	5.72	1.58	3.56	2.47	5.63	2.76			
	NT2RP3003437	3.79	2.73	8.27	7.3	2.16	3.14	2.83	3.31	4.36			
	NT2RP3003448	50.3	43.28	42.61	71.29	41.3	73.4	51.67	39.73	51.96			
	NT2RP3003455	10.05	10.38	13.46	9.15	4.03	7.96	7.2	5.64	6.56	*		-
	NT2RP3003462	10.37	9.93	17.25	9.92	5.42	4.43	5.52	5.56	5.82	*		-
15	NT2RP3003464	14.66	13.89	13.26	12.87	7.49	6.91	18.26	16.54	18.68	**	+	-
	NT2RP3003469	17.22	14.94	13.68	14.5	4.68	9.94	19.05	18.28	19.4	*	+	-
	NT2RP3003473	10.99	11.83	10.53	13.09	12.46	13.39	10.9	7.59	8.71	*		-
	NT2RP3003474	3.37	3.95	4.83	6.73	2.05	5.69	2.42	1.56	2.61	*		-
	NT2RP3003475	25.58	23.74	27.15	19.96	21.31	0.35	11.64	10.85	19.52	*		-
	NT2RP3003490	6.09	2.59	4.63	5.62	1.54	3.26	2.21	16.1	3.36			
20	NT2RP3003491	18.8	17.56	17.3	22.58	14.42	27.63	10.64	13.18	15.87	*		-
	NT2RP3003493	4.17	0.67	1.27	6.14	0.62	2.97	3.62	2.46	1.88			
	NT2RP3003500	5.9	4.89	5.02	7.76	1.8	6.18	6.6	5.32	6.99			
	NT2RP3003527	7.87	4.85	6.47	6.19	2.35	4.17	6.12	6.18	4.88			
	NT2RP3003532	10.61	16.22	14.72	12.47	4.81	8.89	14.42	12.45	12.16			
	NT2RP3003535	38.7	29.85	32.07	25.66	21.86	25.49	14.16	14.9	18.19	*	**	-
25	NT2RP3003536	5.83	2.22	11.18	6.74	1.99	4.05	4.59	4.1	3.31			
	NT2RP3003543	83.01	76.4	71.54	99.52	61.99	69.1	35.13	38.35	47.54	**		-
	NT2RP3003549	5.05	3.61	9.39	4.68	1.49	3.34	2.95	6.92	3.73			
	NT2RP3003552	5.72	3.03	6.03	6.81	1.98	1.74	4.06	3.11	1.34			
	NT2RP3003555	7.3	5.08	11.06	9.46	2.64	6.2	9.1	6.02	7.32			
	NT2RP3003559	16.08	18.05	20.35	13.82	7.72	7.64	10.69	7.28	10	*	**	-
30	NT2RP3003564	21.9	11.18	19.14	32.64	13.35	24.42	5.9	11.04	13.84			
	NT2RP3003572	4.45	3.71	12.53	6.23	1.91	3.3	2.99	5.76	5.28			
	NT2RP3003576	10.13	4.39	10.89	7.2	4.48	6.35	2.89	3.48	6.99			
	NT2RP3003587	4.99	3.94	10.87	4.59	3.45	2.48	4.61	2.31	2.5			
	NT2RP3003589	8.81	6.29	8.93	5.98	6.78	3.45	8.93	5.16	5.31			
	NT2RP3003592	16.97	13.2	13.23	12.11	9.09	12.9	9.83	6.59	9.69	*		-
35	NT2RP3003593	8.95	15.57	14.21	12.38	7.8	13.05	6.6	6.07	15.06			
	NT2RP3003614	16.19	15.38	25.69	16.13	12.3	18.94	17.5	16.33	15.44			
	NT2RP3003621	23.31	9.78	21.55	19.16	8.76	10.65	10.2	15.44	18.25			
	NT2RP3003625	30.15	13.92	27.9	27.09	17.35	16.31	10.08	25.7	26.09			
	NT2RP3003627	13.85	10.22	17.05	10.88	8	10.73	7.14	16.59	27.58			
40	NT2RP3003636	43.56	19.06	24.94	24.57	26.12	25.77	9.71	16.11	10.76			
	NT2RP3003642	9.55	5.41	10.21	6.42	5.7	4.71	4.31	4.84	5.36			
	NT2RP3003645	11.93	7.13	13.91	10.67	5.31	7.85	4.02	4.25	4.25	*		-
	NT2RP3003648	6.56	6.34	9.24	8.44	4.41	4.52	4.4	2.95	3.73	*		-
	NT2RP3003649	26.09	26.77	29.81	22.55	18.26	23.64	34.18	24.81	32.1	*		-
	NT2RP3003650	12.61	11.46	21.64	12.68	5.46	10.24	6.28	12.27	13.31			
	NT2RP3003656	6.54	4.65	10.33	8.68	1.7	4.51	3.42	4.72	11.15			
45	NT2RP3003659	9.46	5.9	15.28	7.57	5.1	6.1	4.35	6.43	5.32			
	NT2RP3003662	8.5	8.39	10.9	5.37	1.92	5.94	1.35	3.6	1.87	*	**	-
	NT2RP3003664	4.6	5.97	8.27	6.79	3.03	4.25	2.14	1.93	3.82	*		-
	NT2RP3003665	11.75	13.45	13.25	12.32	5.11	5.74	4.75	7.06	5.47	**		-
	NT2RP3003671	70.45	72.23	48.58	69.47	88.76	53.27	45.85	13.94	22.44	*		-
50	NT2RP3003672	12.42	9.87	13.46	13.04	14.86	11.85	11.18	14.12	13.48			
	NT2RP3003673	8.49	4.06	14.72	9.56	3.98	4.91	4.97	6.88	5.68			
	NT2RP3003679	10.33	6.46	20.68	8.71	5.36	8.11	4.9	8.58	8.08			
	NT2RP3003680	13.23	15.82	26.63	9.67	11.59	10.92	6.46	10.89	9.67			
	NT2RP3003686	3.02	3.35	12.87	5.63	1.59	1.88	2.06	1.47	1.07			
	NT2RP3003689	9.23	7.33	11.82	8.17	6.77	3.83	2.32	5.97	3.24	*		-
	NT2RP3003697	6.83	7.26	6.91	7.08	5.61	6.06	2.11	3.71	4.84	*		-
55	NT2RP3003701	9.67	9.01	11.76	7.31	4.84	5.19	3.31	2.62	4.46	*	**	-
	NT2RP3003704	10.73	9.76	12.38	10.7	7.93	10.02	15.67	9.66	12.47			
	NT2RP3003714	6.87	2.51	9.56	7.86	2.41	3.23	1.8	2.83	0.8			

Table 447

	NT2RP3003044	20.51	12.05	18.28	17.95	10.92	17.93	12.47	17.92	24.95			
	NT2RP3003047	318.58	233.21	307.78	283.91	158.8	215.31	289.99	339.63	305.02			
5	NT2RP3003050	6.01	3.67	13.67	6.04	1.27	2.97	2.45	4.04	3.78			
	NT2RP3003053	4.73	4.39	9.42	5.4	1.52	3.2	1.65	2.69	4.02			
	NT2RP3003059	3.4	2.18	5.45	4.75	1.77	1.03	2.8	1.96	1.97			
	NT2RP3003061	3.92	4.63	10.62	7.14	5.71	3.17	3.68	3.67	3.35			
	NT2RP3003068	13.64	16.72	19.09	13.94	11.08	9.11	15.51	9.65	9.79			
	NT2RP3003071	501.47	582.68	529.35	407.82	335.95	447.01	379.51	122.11	170.7	*	*	-
10	NT2RP3003076	11.52	4.82	12.85	9.77	5.93	5.76	7.19	9.4	27.45			
	NT2RP3003078	14.69	14.12	17.86	9.88	7.71	7.1	3.81	7.38	11.24	**	*	-
	NT2RP3003081	14.34	14.21	20.22	13.69	9.02	9.23	12.85	17.91	10.55			
	NT2RP3003090	4.38	6.96	9.84	6.19	3.29	2.43	4.26	4.05	2.94			
	NT2RP3003097	14.02	15.06	14.8	11.66	5.57	6.77	8.05	9.19	6.11	*	**	-
	NT2RP3003098	6.1	7.46	9.23	6.48	4.67	4.01	3.99	4.01	3.93	*		-
15	NT2RP3003101	11.04	12.95	14.52	11.99	6.27	8.65	11.09	9.21	8.98			
	NT2RP3003109	27.77	29.32	24.18	32.49	26.06	31.9	22.67	12.21	14.33	*		-
	NT2RP3003121	14.74	6.75	15.15	12.65	4.66	5.96	4.11	5.09	12.75			
	NT2RP3003133	8.34	7.45	12.11	8.93	5.28	6.18	2.52	4.07	7.32			
	NT2RP3003137	4.53	5.11	11.1	4.41	2.55	1.77	1.57	3.58	2.61			
20	NT2RP3003138	14.49	14.02	31.57	15.47	8.83	7.42	8.07	6.92	9.16			
	NT2RP3003139	17.72	13.28	16.88	11.32	8.97	7.87	7.6	10.95	5.4	*	*	-
	NT2RP3003145	29.81	27.36	41.27	37.92	28.86	28.23	19.57	16.3	17.5	*		-
	NT2RP3003150	3.07	5.2	3.99	5.23	1.85	1.66	3.17	2.46	1.85			
	NT2RP3003157	36.81	28.34	31.25	37.59	20.43	24.91	16.55	12.31	12.99	**		-
	NT2RP3003185	12.88	12.16	22.32	13.29	10.08	11.04	7.56	13.67	13.84			
25	NT2RP3003193	56.79	51.34	61.74	61.27	47.04	47.44	61.96	79.57	80.02			
	NT2RP3003197	14.34	9.96	23.45	8.85	5.2	5.54	6.63	9.8	4.5			
	NT2RP3003203	13.42	14.29	19.75	14.19	7.85	8.04	8.52	12.56	10.13			
	NT2RP3003204	19.08	17.04	30.4	16.33	10.47	7.75	7.19	12.58	11.16			
	NT2RP3003210	6.33	8.19	12.29	9.16	9.84	6.29	6.27	7.36	5.46			
	NT2RP3003212	8.37	8.5	11.56	8.16	5.18	3.57	6.2	5.26	6.14	*		-
	NT2RP3003213	5.67	4.05	5.85	6.17	2.96	2.21	4.78	3.56	3.34			
30	NT2RP3003224	55.68	53.67	56.74	16.3	18.72	16.37	11.98	29.14	26.24	**	**	-
	NT2RP3003226	4.57	3.58	10.67	6.47	2.35	3.36	2.91	2.15	2.81			
	NT2RP3003230	7.29	5.28	14.91	8.17	2.63	3.43	3.72	2.45	3.81			
	NT2RP3003235	79.63	66.15	103.5	95.3	92.56	60.75	55.64	70.39	55.23			
	NT2RP3003242	73.44	66.48	77.54	45.7	33.8	25.86	41.52	43.2	38.85	**	**	-
35	NT2RP3003251	104.1	73.55	126.64	135.43	95.34	99.58	79.05	81.02	71.32			
	NT2RP3003252	10.37	11.27	12.13	9.17	5.92	6.54	5.47	5.61	2.73	*	**	-
	NT2RP3003258	15.2	11.75	13.65	15.47	7.62	6.38	7.6	4.08	5.36	**		-
	NT2RP3003260	20.33	21.23	23.55	13.01	8.75	11.13	14.98	15.72	16.47	**	**	-
	NT2RP3003264	39.02	33.18	49.85	37.17	29.58	23.32	18.19	26.52	25.18	*		-
	NT2RP3003273	4.51	4.91	15.07	7.81	4.36	2.59	2.42	2.45	2.89			
40	NT2RP3003278	10.07	13.81	15.32	10.69	5.74	4.93	4.88	5.26	3.08	**		-
	NT2RP3003280	8.01	8.36	9.74	5.35	2.41	1.56	1.8	2.02	2.23	*	**	-
	NT2RP3003282	6.34	4.14	6.3	6.51	1.72	3.27	3.2	1.42	0.91	*		-
	NT2RP3003290	5.35	4.64	5.82	6.64	2.67	3.3	2.13	0.74	2.31	**		-
	NT2RP3003301	10.24	13.13	10.58	8.04	5.22	4.9	4.68	1.19	1.93	*	**	-
	NT2RP3003302	44.6	31.52	44.49	25.69	16.4	28.65	36.54	47.16	60.3	*		-
45	NT2RP3003311	4.75	4.55	11.5	8.43	3.14	4.46	2.2	3.71	1.64			
	NT2RP3003312	24.36	35.16	29.14	9.01	9.9	12.13	5.11	5.48	5.49	**	**	-
	NT2RP3003313	11.85	10.76	12.83	7.95	5.85	3.91	3.77	4.31	3.7	*	**	-
	NT2RP3003327	9.48	7.49	11.81	7.32	3.83	2.89	4.61	4.57	4.3	*		-
	NT2RP3003330	28.77	45.14	40.86	24.02	22.59	18.95	15.05	12.99	8.77	*	**	-
	NT2RP3003340	7.42	9.51	10.17	9.56	3.26	4.53	3.31	3.26	4.42	**		-
50	NT2RP3003346	7.58	7.33	10.55	7.96	3.11	4.95	5.04	6.17	6.71			
	NT2RP3003349	77.06	82.34	60.61	59.72	39.01	41.31	20.2	47.64	67.66	*		-
	NT2RP3003353	5.79	9.35	12.49	5.84	3.66	3.54	2.78	3.58	3.64	*		-
	NT2RP3003356	50.09	93.38	122.44	53.2	56.67	30.98	100.88	61.77	52.32			
	NT2RP3003368	10.01	14.26	18.12	8.66	5.26	5.87	4.9	3.64	2.35	*	*	-
	NT2RP3003375	13.81	11.33	19.09	11.92	5.85	11.47	12.29	9.6	11.66			
55	NT2RP3003377	8.13	11.7	14	8.42	2.73	1.97	7.35	4.71	8.09	*		-
	NT2RP3003384	15.97	18.59	24.94	10.9	5.8	5.57	11.93	4.66	4.64	*	*	-
	NT2RP3003385	10.44	16.16	15.37	14.39	15.35	10.89	10.54	6.35	9.2			

Table 446

	NT2RP3002701	11.4	8.8	15.19	9.64	4.21	7.04	4.79	10.11	7.18			
	NT2RP3002705	30.23	29.89	43.68	30.05	23.71	23.42	20.15	25.03	27.52			
5	NT2RP3002708	25.52	23.92	35.27	19.02	13.81	19.8	19.34	22.92	17.32			
	NT2RP3002711	87.34	128.24	131.86	149.35	126.88	84.65	16.56	37.78	36.54	*	*	-
	NT2RP3002712	11.51	9.81	14.14	7.18	5.15	2.41	5.29	7.06	4.76	*	*	-
	NT2RP3002713	17.13	13.75	17.42	8.41	9.55	4.36	13.41	9.28	8.38	*	*	-
	NT2RP3002721	70.06	117.72	122.3	101.88	70.22	94.12	113.7	122.46	107.46			
	NT2RP3002722	20.45	20.54	15.56	16.97	11.11	10.65	14.55	10.8	10.74	*	*	-
10	NT2RP3002723	12.21	14.76	21.14	7.75	4.55	7.76	4.96	7.72	6.61	*	*	-
	NT2RP3002737	4.67	4.9	12.9	7.24	2.22	4.66	2.62	1.49	3.15			
	NT2RP3002738	122.73	88.31	164.33	156.37	134.11	152.93	81.07	85.63	82.42			
	NT2RP3002742	4.62	5.31	8.06	6.59	2.59	2.11	2.29	3.26	2.78	*		-
	NT2RP3002744	11.76	7.9	17.24	10.03	4.45	6.38	6.88	8.18	6.87			
	NT2RP3002756	8.91	6.64	6.82	7.81	5.13	2.63	2.84	1.31	1.38	**		-
15	NT2RP3002757	18.22	14.86	15.65	13.62	6.41	8.83	10.99	9.29	10.38	*	*	-
	NT2RP3002758	16.5	15.69	16.85	13.98	8.17	9.33	15.83	11.78	10.47	*		-
	NT2RP3002762	6.55	8.17	14.93	9.33	2.38	7.36	4.56	5.11	6			
	NT2RP3002763	6.01	4.94	14.86	7.96	3.26	4.46	4.15	2.77	3.35			
	NT2RP3002770	13.79	15.86	20.19	13.26	7.67	4.65	6.38	6.79	5.16	**		-
	NT2RP3002771	6.48	3.94	12.36	8.16	4.01	1.77	3.51	4.07	2.45			
20	NT2RP3002785	7.93	7.61	12.09	7.52	4.23	4.31	4.14	3.39	4.41	*		-
	NT2RP3002790	12.74	8.68	16.54	10.03	4.49	9.67	11.67	8.3	11.43			
	NT2RP3002799	44.32	36.15	61.61	71.32	40.17	61.97	38.77	29.46	32.11			
	NT2RP3002801	9.13	7.96	8.02	8.82	3.17	7.55	6.99	2.43	2.63	*		-
	NT2RP3002802	6.6	5.16	13.06	9.48	2.83	4.69	3.71	3.1	3.55			
25	NT2RP3002810	21.61	13.43	30.89	15.94	9.02	13.72	15.57	16.49	23.18			
	NT2RP3002818	12.26	21.33	22.29	12.03	10.84	9.9	12.06	10.85	7.02			
	NT2RP3002821	3.08	2.76	7.5	5.13	0.98	2.74	3.88	1.7	1.96			
	NT2RP3002823	5.94	5.9	9.87	7.27	1.84	2.08	2.79	3.12	3.09	*		-
	NT2RP3002825	46.16	68.1	69.32	16.84	16.8	11.22	9.52	7.53	4.62	**	*	-
	NT2RP3002829	36.39	46.01	39.22	22.92	16.02	11.21	24.02	7.98	9.64	**	*	-
30	NT2RP3002831	9.11	9.95	10.41	14.39	9.89	8.12	9.96	4.98	9.9			
	NT2RP3002836	16.68	11.51	17.27	10.91	4.09	9.65	4.06	4.72	7.36	**		-
	NT2RP3002845	38.76	32.47	30.9	12.04	7.1	5.17	5.18	12.27	9.29	**	*	-
	NT2RP3002852	27.07	21.18	24.92	24.24	10.22	18.18	20.47	22.49	18.97			
	NT2RP3002861	6.45	10.44	9.71	7.81	2.41	4.65	6.77	4.65	5.14			
	NT2RP3002869	6.19	9.1	10.7	6.8	3.23	3.7	4.89	3.27	3.99	*		-
35	NT2RP3002874	12.63	13.11	17.2	9.07	5.37	6.1	8.08	3.93	8.06	*	*	-
	NT2RP3002876	143.63	148.27	181.51	254.93	163.87	216.77	137.83	99.6	141.63			
	NT2RP3002877	16.65	19.83	22.97	18.33	13.19	15.19	13.02	8.47	12.29	*		-
	NT2RP3002887	11.91	8.65	14.58	12.86	6.78	9.6	8.16	11.65	6.83			
	NT2RP3002900	24.9	40	47.49	50.72	36.64	57.77	42.16	46.86	41.82			
	NT2RP3002902	50.68	50.65	63.54	17.78	28.94	15.05	19.17	20.83	23.73	**	*	-
40	NT2RP3002909	5.43	3.6	6.73	6.91	1.07	4.91	4.39	3.43	4.07			
	NT2RP3002911	32.27	38.37	39.37	37.75	22.31	32.66	41.96	41.35	47.38			
	NT2RP3002948	97.24	122.54	134.73	183.88	103.17	110.98	109.55	85.09	95.2			
	NT2RP3002953	130.69	155.36	174.07	145.08	99.42	98.1	201.22	181.14	165.58			
	NT2RP3002955	11	13.27	12.07	9.74	5.71	6.63	7.47	5.75	5.17	*	*	-
	NT2RP3002958	11.7	7.18	10.81	11.8	6.8	11.76	8.8	13.15	14.41			
45	NT2RP3002969	10.47	5.67	11.5	11.46	3.57	3.82	2.62	3.28	4.09	*		-
	NT2RP3002972	8.18	3.34	9.31	7.48	3.85	0.87	5.21	5.53	7.19			
	NT2RP3002978	17.8	13.14	17.07	16.61	7.47	16.55	15.56	17.59	21.66			
	NT2RP3002983	14.39	16.02	14.98	15.99	6.52	11.68	18.52	16.38	19.45	*		+
	NT2RP3002985	4.55	5.27	7.26	6.46	1.04	3.38	4.84	3.07	3.43			
	NT2RP3002988	8.52	6.19	9.46	9	4.1	8.27	7.55	8.24	9.04			
50	NT2RP3003000	3.67	3.71	3.5	6.2	1.26	2.48	2.37	2.4	2.73	**		-
	NT2RP3003008	10.96	2.67	10.84	10.43	9.08	7.16	5.31	8.35	5.42			
	NT2RP3003012	2.33	3.47	10.69	4.62	0.4	1.73	0.84	1.69	1.54			
	NT2RP3003015	94.04	42.5	71.87	52.22	37.75	51.79	66.36	44.43	66.31			
	NT2RP3003018	4.14	3.97	9.63	5.98	1.6	1.48	2.56	2.43	2.26			
	NT2RP3003028	3.8	4.39	9.49	7.13	4.25	1.98	3.07	2.68	1.64			
55	NT2RP3003029	10.59	12.64	16.85	9.46	5.29	7.18	11.38	8.45	12.28			
	NT2RP3003032	18.44	16.02	27.77	16.77	11.22	12.42	15.36	17.44	10.46			
	NT2RP3003041	9.55	9.82	9.85	12.39	5.66	4.67	6.47	4.02	3.62	**		-

Table 445

	NT2RP3002377	20.63	15.78	16.8	13.04	9.6	12.99	16.42	23.7	18.15	*	-
	NT2RP3002397	5.32	9.77	13.16	6.81	2.98	3.19	4.39	2.85	3.63		
5	NT2RP3002404	6.54	8.21	7.65	4.33	1.3	1.29	2.68	0.69	1.15	* **	- -
	NT2RP3002410	7.62	4.7	8.02	3.73	3.33	1.2	3.81	3.45	2.67	* *	- -
	NT2RP3002414	5.43	4.93	6.03	6.64	1.85	2.91	4.09	1.68	3.33	*	- -
	NT2RP3002430	5.86	5.44	9.44	4.78	1.52	1.85	3.2	1.23	1.16	*	- -
	NT2RP3002448	12.97	9.25	9.52	5.21	1.95	1.81	2.67	2	4.7	** **	- -
	NT2RP3002454	12.06	19.92	16.37	13.74	7.64	12.02	13.75	9.17	8.67		
10	NT2RP3002455	4.76	3.53	5.11	6.77	1.51	4.71	3.7	5.48	5.26		
	NT2RP3002456	58.03	38.92	43.81	27.96	14.69	15.06	14.57	36.86	33.26	*	-
	NT2RP3002462	20.18	13.39	21.5	10.24	9.33	4.6	3.9	5.28	3.19	* **	- -
	NT2RP3002469	3.83	1.84	6.7	6.03	3.37	1.62	2.45	1.7	0.82		
	NT2RP3002470	35.44	35.5	72.47	34.81	21.72	19.39	33.8	20.85	19.08		
	NT2RP3002484	3.62	2.13	4.4	4.15	1.03	1.2	2.28	1.6	1.75		
15	NT2RP3002491	13.97	21.34	15.21	12.05	5.93	4.04	5.95	7.05	7.23	* *	- -
	NT2RP3002494	24.09	40.78	35.02	15.6	8.77	12.21	26.54	16.61	15.03	*	-
	NT2RP3002497	1.45	2.72	2.12	3.96	0.95	2.47	1.03	1.64	2.67		
	NT2RP3002500	16.58	14.89	20.26	24.82	15.49	27.1	14.57	11.42	19.8		
	NT2RP3002501	32.52	29.62	45.43	41.64	21.97	31.25	12.71	18.89	18.63	*	-
	NT2RP3002512	5.61	3.7	5.13	8.12	4.35	5.53	3.81	5.5	5.08		
20	NT2RP3002529	570.71	676.56	723.55	824.94	401.81	698.08	438.95	899.52	707.21		
	NT2RP3002533	33.09	38.94	36.28	44.79	15.69	31.1	9.54	21.79	17.75	**	-
	NT2RP3002539	6.02	8.36	9.74	6.07	3.74	0.01	2.47	0	0.12	**	-
	NT2RP3002540	4.32	7.97	5.52	9.08	2.78	5.47	9.44	4.32	6.18		
	NT2RP3002543	12.62	4.6	12.88	12.84	5.24	6.49	9.1	11.85	10.42		
	NT2RP3002545	2.99	1.16	9.33	6.12	1.46	2.25	2.07	3.55	0.82		
25	NT2RP3002548	6.99	4.28	13.6	7.32	1.84	2.44	4.48	4.56	2.91		
	NT2RP3002552	4.39	5.59	10.04	7.03	1.31	2.23	3.66	3.6	2.29		
	NT2RP3002558	34.53	21.49	27.53	25.41	11.35	21.08	37	39.71	40.69	*	+
	NT2RP3002565	13.64	7.8	14.18	11.1	3.88	9.31	11.41	13.37	16.44		
	NT2RP3002566	7.8	8.76	13.33	16.41	6.98	8.51	6.63	7.76	8.89		
30	NT2RP3002571	8.13	11.03	10.93	9.57	4.89	4.39	4.63	2.67	3.9	**	-
	NT2RP3002572	74.15	45.74	56.76	41.43	36.41	30.38	19.06	45.1	46.35		
	NT2RP3002573	32.62	15.75	29.51	19.71	16.82	14.99	13.13	13.23	21.22		
	NT2RP3002577	47.28	29.36	41.83	19.87	14.5	21	17.59	15.85	17.3	* *	- -
	NT2RP3002579	8.63	6.06	12.6	7.23	3.12	4.38	5.33	4.66	5.1		
	NT2RP3002582	6.19	3.82	10.45	7.84	3.56	3.38	4.87	4.78	3.74		
	NT2RP3002587	10.07	10.45	15.76	10.84	6.77	7.46	4.8	4.74	3.88	*	-
35	NT2RP3002590	544.94	699.14	857.26	489.72	627.82	453.24	363.97	273.66	582.87		
	NT2RP3002602	7.18	7.63	10.58	10.16	7.38	6.68	4.87	4.34	6.16	*	-
	NT2RP3002603	14.44	13.52	18.75	16.38	10.66	15.03	8.16	15.4	88.12		
	NT2RP3002621	6.81	5.15	9.83	8.56	3.04	3.97	3.12	5.17	6.09		
	NT2RP3002622	6.75	5.01	13.82	8.8	3.27	3.73	2.74	4.44	11.45		
40	NT2RP3002624	26.96	16.99	21.5	17.24	10.36	13.21	17.97	25.27	18.08		
	NT2RP3002628	6.59	5.38	9.7	6.43	2.65	4.46	4.24	5.21	2.97		
	NT2RP3002629	5.76	11.73	14.59	10	6.43	5.93	6.71	4.48	3.96		
	NT2RP3002631	48.35	45.23	46.18	26.1	22.1	27.75	26.27	23.38	21.25	** **	- -
	NT2RP3002647	12	15.89	13.32	16.2	15.29	15.81	10.24	7.5	7.23	*	-
	NT2RP3002649	9.3	7.92	12.59	9.96	4.23	5.01	3.45	5.1	5.57	*	-
45	NT2RP3002650	26.86	18.53	27.47	18.25	9.68	13.45	17.35	20.46	18.82		
	NT2RP3002652	5.52	5.57	14.03	5.61	3.76	3.01	2.75	4.84	3.27		
	NT2RP3002654	20.14	13.49	14.7	11.81	10.11	6.69	5.42	7.8	5.1	*	-
	NT2RP3002657	16.5	15.29	20.85	14.05	6.31	13.07	15.2	14.6	14.98		
	NT2RP3002659	15.92	19.34	18.66	13.66	10.7	8.74	4.85	10.09	6.73	* **	- -
	NT2RP3002660	5.66	10.19	9.81	8.43	3.68	4.95	6.72	4.39	3.92		
50	NT2RP3002663	10.53	17.57	11.68	14.14	15.43	9.63	19.1	11.04	8.75		
	NT2RP3002664	28.81	23.63	39.16	18.72	11.43	13.27	7.92	17.21	15.51	* *	- -
	NT2RP3002687	32.83	27.15	33.62	23.94	18.84	19.81	15.18	21.3	26.19	*	-
	NT2RP3002671	38.88	30.21	40.39	24.11	16.4	14.74	30.8	35.93	33.74	*	-
	NT2RP3002682	8.45	6.51	13.68	7.64	3.93	4.02	2.41	3.04	2.62	*	-
	NT2RP3002684	20.03	14.16	18.81	8.9	9.81	5.51	4.13	6.96	4.79	* **	- -
55	NT2RP3002687	9.25	11.21	13.71	14.12	8.33	10.78	7.86	7.74	7.01	*	-
	NT2RP3002688	168.35	106.47	188.15	190.36	164.88	199.08	153.97	87.55	117.43		
	NT2RP3002698	8.07	8.3	9.14	8.34	5.8	6.56	5.59	5.87	4.73	**	-

Table 444

	NT2RP3002086	9.1	5.64	14.68	7.72	4.13	4.75	9.35	9.69	7.69			
	NT2RP3002094	3.34	2.83	6.44	4.61	2.06	2.3	2.41	5.92	1.29			
5	NT2RP3002096	12.33	8.92	11.73	8.59	3.97	5.5	5.94	5.18	4.01	*	**	-
	NT2RP3002097	23.91	20.85	34.1	19.75	11.2	19.57	12.95	19.73	14.4			
	NT2RP3002098	5.03	5.84	6.49	7.72	2.61	2.62	5.48	4.78	3.63			
	NT2RP3002102	13.2	5.06	15.39	11.41	5.66	7.42	3.96	4.11	3.36			
	NT2RP3002106	10.36	7.3	14.64	10.63	8.57	6.26	3.35	4.87	10.6			
	NT2RP3002108	4.96	2.64	12.49	5.65	1.44	2.35	2.54	2.99	5.43			
10	NT2RP3002109	40.98	15.02	38.55	15.39	16.89	11.31	8.11	16.82	8.14			
	NT2RP3002110	5.34	2.87	6.32	5.23	2.17	1.37	2.33	2.43	0.38			
	NT2RP3002113	9.48	7.75	15.49	10.55	5.09	3.01	6.14	4.67	6.08			
	NT2RP3002120	7.73	8.74	11.8	6.57	3.91	2.89	5.04	3.74	4.99	*	*	-
	NT2RP3002121	1.11	4.94	1.77	5.17	1.32	0.68	2.8	1.37	1.2			
	NT2RP3002126	64.09	54.26	55.37	40.43	28.8	44.09	33.92	65.97	39.23	*		-
15	NT2RP3002128	8.94	4.6	9.8	7.18	4.51	3.23	4.1	5.32	11.38			
	NT2RP3002130	45.15	21.81	38.29	25.97	19.96	17.62	5.74	23.29	23.75			
	NT2RP3002133	713.78	729.07	666.05	842.53	766.78	697.72	544.62	1065.2	813.76			
	NT2RP3002136	99.29	50.26	79.88	69.55	58.35	39.45	17.4	37.71	52.59			
	NT2RP3002140	3.69	7.52	7.94	5.4	2.45	3.08	2.99	3.81	3.42			
	NT2RP3002142	9.07	12.94	11.89	8.33	6.6	5.19	5.4	5.06	2.64	*	**	-
20	NT2RP3002146	9.3	10.61	11.18	9.37	7.32	7.23	12.19	11.03	8.72			
	NT2RP3002147	11.07	11.26	19.71	10.63	7.44	5.02	4.38	6.88	4.67	*		-
	NT2RP3002151	15	10.97	25.17	14.01	7.18	7.82	7.02	15.09	18.04			
	NT2RP3002155	53.25	66.04	47.55	73.17	43.36	52.61	36.35	94.67	86.47			
	NT2RP3002156	3.79	2.7	6.74	4.59	0.75	1.4	1.04	0.59	1.13	*		-
	NT2RP3002160	9.08	10.25	12.47	12.87	7.96	7.04	2.86	3.65	6.87	*		-
25	NT2RP3002163	13.64	21.03	16.3	10.69	7.06	4.19	9.73	11.32	12.78	*		-
	NT2RP3002165	6.75	7.58	8.41	6.08	1.96	1.96	2.6	3.84	1.9	*	**	-
	NT2RP3002166	5.43	5.03	4.69	5.27	1.55	0.67	2.88	6.31	2.47			
	NT2RP3002173	517.9	446.69	509.69	297.97	240.6	253.29	44.66	135.22	128.7	**	**	-
	NT2RP3002174	130.22	187.95	156.75	150.48	138.99	121.68	76.48	86.96	88.77	*		-
30	NT2RP3002181	5.97	4.69	12.81	5.8	2.39	3.23	1.84	5.23	1.34			
	NT2RP3002185	44.31	38.54	51.77	25.79	16.42	22.4	38.06	52.39	48.7	**		-
	NT2RP3002193	7.33	7.63	13.87	8.26	3.71	3.99	4.19	5.78	8.23			
	NT2RP3002204	41.05	45.66	52.03	45.02	28.36	25.63	28.31	47.28	47.76			
	NT2RP3002244	12.8	20.63	18.31	13.33	15.24	10.05	9.17	7.08	5.13	*		-
	NT2RP3002248	7.98	7.8	12.04	10.57	4.97	3.5	9.37	6.63	3.98			
35	NT2RP3002253	17.21	11.16	18.79	9.56	7.33	7.05	3.72	10.84	9.39	*		-
	NT2RP3002255	3.26	2.63	8.57	4.41	1.55	2.02	1.36	3.59	1.93			
	NT2RP3002264	81.68	74.91	64.3	98.72	70.97	43.12	54.37	98.78	53.26			
	NT2RP3002267	2.25	3.31	8.07	3.82	0.82	1.41	0.79	1.35	0.88			
	NT2RP3002273	20.4	16.49	19.5	15.02	8.97	5.71	10.11	9.27	13.28	*	**	-
	NT2RP3002276	3.97	2.93	8.36	4.35	0.7	1.55	1.24	0.89	0.84			
40	NT2RP3002281	13.35	12.31	15.77	11.45	7.06	7.28	10.16	10.76	11.32	*	*	-
	NT2RP3002286	16.62	19.5	20.21	12.28	6.1	7.69	4.25	3.16	1.57	**	**	-
	NT2RP3002297	94.9	98.81	132.4	64.72	33.84	39.88	11.41	37.45	29.88	*	**	-
	NT2RP3002301	11.34	12.34	18.57	7.11	3.98	4.21	2.56	4.64	4.68	*	*	-
	NT2RP3002303	7.54	10.15	16.1	7.32	5.62	4.17	5.05	6.82	7.85			
	NT2RP3002304	91.67	49.25	50.79	81	77.65	34.95	31.29	39.07	25.04			
45	NT2RP3002309	4.32	3.35	10.38	4.01	1.44	1.07	2.59	1.36	0.22			
	NT2RP3002311	9.65	9.45	12.73	9.08	6.57	4.41	3.9	4.09	5.31	**		-
	NT2RP3002315	16.31	24.1	16.79	7.96	7.42	8.29	9.67	8.19	10.65	*	*	-
	NT2RP3002319	8.49	14.45	9.47	9.99	3.2	4.43	5.43	3.04	5.3	*		-
	NT2RP3002324	18.48	20.79	21.07	16.08	9.05	11.92	2.65	9.16	8.78	*	**	-
	NT2RP3002330	14.44	26.39	29.36	13.92	12.3	11.42	5.88	11.72	10.72	*		-
50	NT2RP3002333	41.51	67.04	49.57	36.81	41.81	35.64	33.68	44.76	37.87			
	NT2RP3002337	4.44	8.96	11.93	5.8	1.82	3.91	2.32	2.11	2.55	*		-
	NT2RP3002342	25.61	22.15	24.13	19.61	11.87	9.75	11.85	13.94	20.4	*	*	-
	NT2RP3002343	3.99	7.39	7.92	5.18	2.92	2.42	2.4	1.77	3.64	*		-
	NT2RP3002351	20.96	17.51	19.58	6.79	4.97	3.6	9.93	4.83	3.54	**	**	-
	NT2RP3002352	29.92	68.54	51.75	25.45	36.1	22.42	47.8	10.88	7.27			
55	NT2RP3002353	26.94	28.46	26.72	23.44	18.05	15.48	14.2	29.02	22.52	*		-
	NT2RP3002362	18.42	12.38	24.27	11.02	6.28	7.66	12.53	15.38	16.92			
	NT2RP3002363	11.71	14.02	21.35	9.9	11.24	7.29	10.05	11.93	6.97			

Table 443

	NT2RP3001764	3.84	4.67	12.85	4.07	1.87	1.79	1.19	3.02	2.08			
	NT2RP3001771	10.53	6.71	15.52	12	8.69	4.55	2.42	4.06	4.07	*		-
5	NT2RP3001777	40.65	47.79	33.95	47.7	28.57	21.63	29.91	18.82	40.09			
	NT2RP3001782	18.47	18.59	14.66	13.14	8.87	10.49	5.89	13.23	9.56	*	*	-
	NT2RP3001792	51.98	43.73	63.18	42.18	33.54	33.35	80.48	61.45	65.58	*		-
	NT2RP3001799	7.5	9.23	9.91	6.52	4.07	3.35	7.42	5.61	3.54	*		-
	NT2RP3001819	9.21	8.05	11.94	7.62	3.17	5.23	2.57	7.27	6.12			
	NT2RP3001829	0.79	0.98	6.7	3.6	0.17	1.25	3.26	1.13	0.11			
10	NT2RP3001836	16.84	18.26	30.32	14.13	7.32	15.63	6.9	7.75	7.46	*		-
	NT2RP3001839	6.81	9.19	13.79	8.6	4.4	4.34	2.3	5.79	3.54			
	NT2RP3001844	5.94	4.03	11.89	5.39	1.37	2.09	2.04	1.24	1.53			
	NT2RP3001848	53.13	36.02	38.27	40.15	39.26	28.02	46.65	51.58	51.5			
	NT2RP3001854	53.05	46.79	83.54	47.02	38.11	28.76	72.37	61.23	74.82			
	NT2RP3001859	8.77	9.33	11.43	9.53	8.22	8.62	9.1	7.34	5.15			
15	NT2RP3001857	2.78	1.95	7.57	5.8	0.8	2.63	1.53	2.14	0.96			
	NT2RP3001858	1.48	1.77	7.55	6.01	1.18	1.83	0	1.41	1.37			
	NT2RP3001861	4.86	3.76	12.83	6.03	2.09	1.54	1.75	2.42	0.53			
	NT2RP3001866	4.29	4.19	8.56	7.38	2.36	3.86	2.27	2.97	2.71			
	NT2RP3001871	4.11	2.37	11.14	4.75	0.71	0.4	0.62	1.36	0.06			
20	NT2RP3001874	10.44	5.38	11.3	5.07	2.46	1.93	2.08	3.65	5.3	*		-
	NT2RP3001878	6.84	8.82	7.25	6.09	2.26	2.81	4.01	2.14	3.46	*	**	-
	NT2RP3001885	6.75	5.91	9.28	7.33	5.82	5.26	4.29	3.86	3.96	*		-
	NT2RP3001896	15.8	9.61	18.78	10.19	5.77	12.23	2.88	7.21	8.15			
	NT2RP3001898	10.12	6.81	17.59	9.29	3.83	4.48	2.35	4.36	3.15			
	NT2RP3001899	4.46	3.48	9.66	6.74	4	4.77	3.33	4.46	3.02			
	NT2RP3001901	2.97	2.49	8.61	4.89	2.11	2.05	1.6	0.22	0.74			
25	NT2RP3001915	21.98	18.4	24.23	18.83	15.46	12.75	6.48	6.95	5.98	**		-
	NT2RP3001926	42.5	70.22	59.74	46.58	37.18	35.96	45.08	32.42	25.88			
	NT2RP3001929	7.68	7.59	9	5.33	1.89	5.28	3.26	1.47	1.63	*	**	-
	NT2RP3001931	7.03	5.58	8.39	8.57	1.83	4.32	4.68	3.69	4.06	*		-
	NT2RP3001938	0.69	0.54	7.42	4.87	0	1.52	0	1.16	0.26			
30	NT2RP3001943	188.81	183.6	152.87	80.5	86.76	128.25	85.31	121.73	115.26	*	*	-
	NT2RP3001944	17.12	20.99	26.58	14.69	15.73	4.63	10.12	15.18	9.75	*		-
	NT2RP3001945	39.73	40.32	37.82	24.74	26.95	28.23	17.61	14.81	14.26	**	*	-
	NT2RP3001947	47.48	38.36	48.19	30.55	15.52	22.25	24.69	18.23	25.96	*	**	-
	NT2RP3001949	37.29	38.85	47.2	29.25	26.76	28.05	30.76	28.58	28.74	*	*	-
	NT2RP3001952	10.17	10.07	6.99	3.68	1.57	0.37	2.44	1.18	1.17	**	**	-
35	NT2RP3001954	9.56	18.54	17.64	8.37	4.32	7.01	11.86	5.36	4.75	*		-
	NT2RP3001956	3.88	4.98	8.37	7.17	1.4	2.11	0.35	2.86	1.91			
	NT2RP3001967	7.48	3.68	9.02	8.89	3.53	3.86	3.05	4.36	2.97			
	NT2RP3001968	2.15	1.05	4.78	4.72	1.42	2.2	1.63	2.56	0.82			
	NT2RP3001976	1.89	2.22	5.25	5.93	1.38	2.82	1.33	1.09	1.87			
	NT2RP3001986	3.89	3.3	6.39	4.22	1.46	2.65	3.43	6.44	2.42			
40	NT2RP3001989	8.56	8.64	5.36	8.05	1.81	1.73	3.37	1.97	1.12	*		-
	NT2RP3002002	7.76	11.38	8.57	5.38	2.01	2.22	6.25	3.04	6.16	*		-
	NT2RP3002004	7.76	12.41	12.06	6.85	3.16	4.05	3.73	2.48	3.6	*	**	-
	NT2RP3002007	14.2	10.34	19.96	11.2	4.92	6.67	4.6	9.34	6.75			
	NT2RP3002014	2.11	1.84	5.64	7.64	3.81	2.33	1.74	3.76	3.1			
	NT2RP3002015	10.24	10.71	16.79	11.66	11.36	10.59	8.67	8.67	11.36			
45	NT2RP3002033	8	5.97	11.92	6.98	3.29	5.92	6.91	8.34	9.48			
	NT2RP3002045	7.45	16.7	16.88	5.97	3.53	4.5	4.78	3.11	3.05	*	*	-
	NT2RP3002054	292.36	495.82	456.61	206.09	224.87	171.66	441.79	291.3	543.12	*		-
	NT2RP3002056	3.09	3.6	0.88	3.45	0.32	0.79	0.92	1.02	0.67			
	NT2RP3002057	10.28	15.21	13.61	7.78	1.9	3.69	8.33	6.11	3.21	*	*	-
	NT2RP3002061	3.7	2.82	3.37	5.14	1.22	2.6	0	1.37	0.53	**		-
50	NT2RP3002062	14.95	11.22	12.96	12.28	11.33	13.44	7.88	11.32	11.12			
	NT2RP3002063	3.07	0.91	3.18	5.76	3.45	2.53	1.35	2.57	1.94			
	NT2RP3002064	4.75	4.07	6.08	8.44	3.11	3.26	2.69	6.61	3.66			
	NT2RP3002071	2.99	2.02	2.84	4.58	1.41	2.26	2.09	2.02	0.69			
	NT2RP3002073	4.16	1.73	3.01	5.39	0.35	0	3.4	1.98	3.29			
	NT2RP3002074	3.97	4.7	9.59	5.64	1.52	1.74	2.26	2.79	1.06			
55	NT2RP3002075	115.61	136.72	152.03	61.15	60.41	66.84	138.08	127.44	140.01	**		-
	NT2RP3002077	1.88	1.37	7.12	4.73	0	2.99	1.52	0.95	1.4			
	NT2RP3002081	9.52	4.64	17.43	9.04	2.14	3.36	6.77	6.6	8.59			

Table 442

	NT2RP3001479	6.51	3.77	10.27	6.45	4.18	2.2	2.6	4.56	5.3			
	NT2RP3001490	15.72	6.25	23.45	15.28	5.31	6.47	7.24	7.84	8.04			
5	NT2RP3001492	14	14.91	21.99	13.76	7.53	9.81	14.28	13.21	12.48			
	NT2RP3001495	18.56	21.11	19.03	12.08	9.76	9.93	15.89	18.33	21.35	**	-	
	NT2RP3001497	36.57	27.99	40.37	47.7	28.75	36.07	23.59	21.43	24.3	*	*	-
	NT2RP3001501	16.76	28.67	29.3	12.62	8.54	7.31	11.78	2.93	8.97	*	*	-
	NT2RP3001527	5.54	5.56	5.02	7.34	2.65	3.31	4.71	3	2.59	*	*	-
	NT2RP3001529	21.9	27.01	22.29	20.64	12.59	16.78	7.77	18.39	18.96			
10	NT2RP3001538	80.06	67.26	76.24	89.51	76.4	114.47	59.99	77.08	72.65			
	NT2RP3001539	4.24	4.99	8.09	6.1	3.34	4.14	3.58	5.72	3.56			
	NT2RP3001542	17.53	25.89	26.44	14.74	14.15	8.68	14.26	12.41	17.97	*	-	
	NT2RP3001549	6.14	6.2	5.77	5.79	1.73	1.92	2.67	2.05	1.89	*	*	-
	NT2RP3001554	28.4	40.66	43.76	10.47	10.16	7.96	15.92	6.96	13.22	**	**	-
	NT2RP3001560	79.18	91.03	86.96	84.47	48.32	76.97	115.31	93.32	84			
15	NT2RP3001561	3.47	3.93	2.29	6.28	2.06	3.29	2.23	2.16	2.49			
	NT2RP3001564	34.74	32.97	34.99	17.25	16.65	19.41	10.11	30.58	15.23	**	-	
	NT2RP3001568	23.08	14.44	30.67	16.92	14.73	6.11	7.97	12.05	7.85			
	NT2RP3001575	34.43	18.91	39.8	19.69	12.66	21.53	16.47	22.21	25.65			
	NT2RP3001580	17.53	21.3	27.37	11.9	7.8	11.25	7.5	7.41	8.67	*	*	-
	NT2RP3001587	14.32	16.9	20.18	11.06	6.6	9.89	19.28	18.65	20.31	*	-	
20	NT2RP3001589	6.31	5.44	5.39	5.29	2.4	2.48	2.34	4.87	3.41			
	NT2RP3001592	37.3	61.22	57.18	30.41	23.17	28.18	46.3	38.66	32.65	*	-	
	NT2RP3001607	11.49	11.68	9.02	7.34	3.49	8.45	5.15	2.53	3.96	**	-	
	NT2RP3001608	30.61	24.16	28.98	15.14	18	18.45	9.66	10.92	10.91	**	**	-
	NT2RP3001613	55.34	46.31	54.71	81.86	48.17	86.77	51.11	48.11	44.78			
25	NT2RP3001619	162.64	181.38	185.09	152.03	118.43	125.85	144.48	167.59	158.06	*	-	
	NT2RP3001621	29.97	24.25	25.08	33.83	16.83	30.34	32.22	27.66	24.11			
	NT2RP3001629	28	13.33	26.21	16.28	12.59	19.85	19.79	23.48	12.56			
	NT2RP3001630	3.51	2.38	4.56	4.63	1.01	2.11	3.79	2.64	3.67			
	NT2RP3001631	64.16	51.12	67.22	112.91	65.5	60.67	59.38	50.82	71.84			
	NT2RP3001634	3.71	4.65	2.94	7.85	2.25	2.09	4.62	2.51	1.18			
30	NT2RP3001642	29.24	22.78	34.25	25.21	18.32	21.46	34.78	35.92	40.14			
	NT2RP3001646	18.72	13.31	23.84	17.34	7.46	8.49	11.13	12.12	12.05			
	NT2RP3001650	4.88	6.33	12.48	7.77	5.89	4.37	3.47	3.76	3.82			
	NT2RP3001667	17.36	19.67	39.76	21.84	7.89	10.07	11.81	14.81	13.16			
	NT2RP3001671	13.32	10.22	13.86	10.05	5.86	9.27	7.9	6.32	3.97	*	-	
	NT2RP3001672	56.18	72.22	87.94	28.72	18.36	21.55	10.7	4.04	14.95	**	**	-
35	NT2RP3001676	11.94	10.66	12.76	8.88	5.89	6.36	7.09	3.31	6.98	*	*	-
	NT2RP3001678	368.4	207.45	268.13	268.3	249.2	237.9	189.86	337.31	274.16			
	NT2RP3001679	19.45	8.41	17.68	13.47	5.77	7.99	3.71	6.54	7.99			
	NT2RP3001682	115.37	82.43	118.94	71.89	66.26	91.02	56.88	114.15	137.5			
	NT2RP3001685	4.91	3.52	11.94	5.16	2.78	2.73	5	2.99	2.91			
	NT2RP3001688	4.75	1.41	4.66	3.33	0.27	0.64	0.99	2.07	0			
40	NT2RP3001690	8.33	10.1	11.83	11.07	4.69	5.02	5.74	5.13	6.32	*	-	
	NT2RP3001693	2.22	6.17	10.5	5.55	2.56	2.65	2.84	2.33	2.39			
	NT2RP3001696	7.34	11.29	11.31	6.93	5.94	11.43	6.97	3.31	6.44			
	NT2RP3001698	780.45	572.31	725.4	616.13	658.96	0.94	533.83	1241.6	1289.4			
	NT2RP3001708	348.43	297.84	257.74	514.57	355.27	1.82	148.2	628.13	519.33			
	NT2RP3001712	32.9	21.37	35.65	18.32	14.72	16.28	22.44	33.16	22.67	*	-	
45	NT2RP3001716	13.86	9.55	15.36	8.27	11.28	8.27	5.55	15.45	9.22			
	NT2RP3001724	28.26	21.7	18.91	32.47	31.93	25.73	16.65	23.86	18.15			
	NT2RP3001727	27.92	15.82	33.03	18.01	12.47	12.24	8.97	9.14	8.89	*	-	
	NT2RP3001729	2.64	5.64	5.12	5.09	2.61	2.56	2.6	2.88	1.65			
	NT2RP3001730	421.81	524.88	490.91	328.42	274.16	316.08	179.32	117.74	101.94	**	**	-
	NT2RP3001733	1.54	1.82	4.9	4.53	0	1.49	0.54	0.39	1.87			
50	NT2RP3001737	7.85	25.52	17.42	10.76	4.14	4.59	3.35	7.04	17.36			
	NT2RP3001738	28.07	24.45	41.36	18.76	13.82	15.18	12.56	15.63	32.84	*	-	
	NT2RP3001739	2.19	2.42	9.16	4.78	1.8	1.62	1.06	2.07	0.9			
	NT2RP3001742	11.1	11.43	14.41	9.77	5.51	6.65	7.73	6.52	8.12	*	*	-
	NT2RP3001751	1.19	1.87	3.32	5.53	0.3	1.67	1.29	0.92	0.82			
	NT2RP3001752	2.48	4.8	2.87	3.47	0.21	1.52	1.75	2.38	0.9			
55	NT2RP3001753	3.96	5.83	5.77	7.82	4.31	2.86	4.73	5.28	2.61			
	NT2RP3001754	3.14	1.28	9.55	3.89	0.47	2.48	0.99	1.82	0.86			
	NT2RP3001756	4.67	4.56	13.42	7.12	2.77	3.58	2.62	4.94	3.56			

Table 441

	NT2RP3001236	30.11	19.17	30.67	17.48	11.33	17.06	7.98	12.61	12.49	*	-
	NT2RP3001239	12.58	7.93	16.69	7.8	5.43	5.92	7.31	5.36	3.8		
5	NT2RP3001240	16.64	15.42	32.69	20.34	13.57	11.75	7.09	11.44	8.12		
	NT2RP3001245	8.04	12.54	22.57	10.74	8.24	10.53	7.27	7.75	6.31		
	NT2RP3001253	5.41	9.93	13.75	5.39	2.74	1.6	4.68	3.73	2.31		
	NT2RP3001259	278.9	310.98	327.25	256.7	262.84	274.28	334.89	289.89	242.91		
	NT2RP3001260	18.94	14.44	22.33	34.17	17.19	23.22	21.96	25.77	20.83		
	NT2RP3001264	67.04	56.89	77.24	82.42	71.22	82.52	43.6	44.18	61.74		
10	NT2RP3001268	54.13	43.25	63.3	45.53	25.93	35.18	39.08	48.29	54.07		
	NT2RP3001271	7.61	5.79	12.88	6.16	2.76	3.69	3.87	7.7	5.38		
	NT2RP3001272	23.18	22.6	31.48	18.93	15.25	15.77	18.56	19.39	20.08	*	-
	NT2RP3001274	143.91	121	164.35	135.41	115.24	154.69	133.08	136.37	139.51		
	NT2RP3001275	7.03	7.19	20.08	7.2	8.45	4.21	4.17	4.56	3.94		
	NT2RP3001280	14.64	18.72	19.46	9.93	11.17	5.67	7.5	8.91	6.46	*	**
15	NT2RP3001281	0.38	2.63	0.7	4.71	1.73	1.17	1.88	0.77	0.6		
	NT2RP3001288	43.86	21.34	42.34	25.62	18.43	26.29	27.01	32.19	37.1		
	NT2RP3001297	77.1	64.75	81.93	35.55	27.23	38.6	12.09	28.24	29.77	**	**
	NT2RP3001300	9.48	9.06	13.81	7.61	5.56	4.66	4.6	5.82	8.28		
	NT2RP3001301	5.4	3.52	7.37	3.86	1.62	1.26	1.66	3.26	1.26		
	NT2RP3001307	13.22	10.36	15.55	10.66	10.08	6.04	5.09	8.17	7.35	*	-
20	NT2RP3001310	9.88	12.87	16.58	11.85	7.95	6.71	4.27	4.42	3.89	*	-
	NT2RP3001318	7.4	8.54	9.21	7.25	6.79	6.73	5.34	5.32	3.43	*	-
	NT2RP3001322	11.09	11.38	14.04	15.94	12.19	11.47	14.66	8.17	7.73		
	NT2RP3001325	1201.7	1025	1099	286.93	377.86	405.84	75.06	206.06	219.63	**	**
	NT2RP3001338	162.73	132.54	198.96	171.74	181.83	241.04	134.27	161.16	143.19		
	NT2RP3001339	16.04	11.7	17.8	11.35	7.84	6.88	2.83	5.64	4.58	*	**
25	NT2RP3001340	35.84	31.42	47.79	30.03	31.99	45.26	31.64	27.19	27.6		
	NT2RP3001341	37.44	66.22	47.17	33.31	24.96	24.34	25.96	38.56	34.37		
	NT2RP3001354	42.85	32.23	43.99	23.31	23.56	19.16	13.01	13.77	11.4	*	**
	NT2RP3001355	2.26	7.76	10.15	5.3	3.13	3.72	4.66	3.96	2.91		
	NT2RP3001356	43	54.12	51.32	62.98	37.84	46.81	81.45	50.61	59.12		
	NT2RP3001359	34.72	29.05	42.17	41.55	33.32	37.07	21.86	23.34	24.66	*	-
30	NT2RP3001364	19.26	26.46	29.33	17.86	11.52	13.89	13.78	18.18	15.95	*	*
	NT2RP3001373	16.79	12.24	26.6	10.58	10.1	9.34	9.17	17.19	12.49		
	NT2RP3001374	95.59	73.12	108.77	136.19	114.69	107.19	68.33	56.48	69.32		
	NT2RP3001383	12	19.08	15.5	12.1	5.75	6.51	13.27	9.62	6.82		
	NT2RP3001384	22.91	17.21	30.7	33.57	22.7	20.56	19.13	14.74	15.12		
	NT2RP3001388	9.38	9.17	7.78	9.06	5.69	4.27	7.73	5.71	5.18	*	-
35	NT2RP3001392	4.64	4.2	2.65	5.43	2.03	2.78	2.37	2.94	3.1		
	NT2RP3001396	47.27	37.6	60.04	23.62	14.23	34.71	12.93	30.54	38.42		
	NT2RP3001398	13.09	9.85	15.48	8.92	6.22	7.1	4.4	8.04	6.53	*	*
	NT2RP3001399	17.91	14.49	24.37	16.05	7.15	6.8	6.07	7.51	6.62	*	-
	NT2RP3001402	30.22	31.51	33.24	60.13	23.85	9.5	15.21	24.77	23.82	*	-
40	NT2RP3001407	72.56	60.86	88.74	96.01	74.06	85.12	43.61	63.39	61.28		
	NT2RP3001416	18.88	22.22	16.48	16.79	17.16	5.73	3.31	4.98	5	**	-
	NT2RP3001420	5.88	3.11	4.79	4.91	2.62	1.74	1.34	1.67	2.52	*	-
	NT2RP3001425	11.74	10.47	11.8	11.91	6.37	9.11	8.4	4.94	6.04	*	-
	NT2RP3001426	12.17	11.79	16.88	9.1	9.64	6.43	5.51	9.62	7.81	*	-
	NT2RP3001427	10.81	11	21.59	12.3	8.43	8.87	17.63	22.78	19.89		
45	NT2RP3001428	7.7	10.58	20.16	8.58	3.38	6.04	5.05	8.84	8.9		
	NT2RP3001429	4.55	6.04	12.09	5.05	1.79	1.93	2.26	1.46	1.89		
	NT2RP3001432	20.32	18.99	30.96	16.39	14.24	12.25	16.49	28.23	11.89		
	NT2RP3001439	31.43	37.32	28.95	26.93	25.44	19.21	14.96	11	10.21	**	-
	NT2RP3001441	34.77	41.28	46.38	53.64	42.72	50.36	30.69	23.25	24.74	*	-
	NT2RP3001446	4.24	2.49	4.94	4.53	3.01	2.2	2.82	2.01	1.51		
50	NT2RP3001447	22.35	21.24	25.11	13.3	8.65	15.94	13.67	37.08	14.99	*	-
	NT2RP3001448	139.79	126.76	203.02	199.4	166.47	178.45	131.32	106.74	139.38		
	NT2RP3001453	31.6	35.5	49.48	34.98	29.3	36.04	28.49	31.86	35.69		
	NT2RP3001457	13.38	17.23	17.13	11.75	10.12	6.15	5.98	6.75	8.92	*	**
	NT2RP3001459	11.29	13.84	13.09	7	3.75	5.3	5.53	4.44	5.41	**	**
	NT2RP3001463	6.65	4.84	3.81	4.67	1.94	1.07	1.96	2.08	0.69	*	-
55	NT2RP3001466	5.28	4.9	5.4	5.77	2.51	1.55	2.18	1.94	1.06	**	-
	NT2RP3001472	5.37	5.26	7.06	12.24	7.01	3.59	1.12	2.87	2.9	*	-
	NT2RP3001475	230.39	301.8	150.57	92.73	106.13	95.77	29.54	68.35	93.78	*	*

Table 440

	NT2RP3000895	2.84	2.51	12.27	4.16	1.17	2.03	1.28	3.4	1.59			
	NT2RP3000900	10.4	8.3	14.31	18.92	3.45	5.38	4.15	13.51	4.93			
5	NT2RP3000901	14.3	18.5	22.78	13.03	13.47	11.77	11.97	9.53	14.53			
	NT2RP3000903	5.01	6.85	6.69	5.82	1.64	3.35	1.28	1.92	1.35	**	-	
	NT2RP3000904	108.6	100.47	123.83	118.07	108.3	100.81	97.74	77.69	108.28			
	NT2RP3000907	84.64	90.19	89.66	84.77	80.01	92.59	125.26	82.21	80.83			
	NT2RP3000913	19.19	19.49	20.62	24.35	9.57	18.44	17.39	14.75	19.21			
10	NT2RP3000917	16.28	18.3	30.01	16.96	15.22	12.11	11.68	16.63	14.88			
	NT2RP3000919	9.33	7.65	14.92	9.52	7.41	6.8	4.46	8.2	5.92			
	NT2RP3000921	112.89	136.59	133.65	119.03	103.41	74.11	25.71	35.79	29.01	**	-	
	NT2RP3000942	67.7	68.21	86.84	75.24	51.25	30.13	38.31	52.76	48.83	*	-	
	NT2RP3000968	50.83	31.86	37.11	29.89	24.92	14.66	9.86	11.81	12.71	**	-	
	NT2RP3000974	8.96	8.74	9.35	9.04	12.66	5.51	4.39	4.92	2.99	**	-	
15	NT2RP3000980	52.51	65.55	64.13	24.76	40.04	30.49	19.39	8.08	16.43	**	**	-
	NT2RP3000984	21.27	24.19	17.57	32.72	12.14	21.58	18.47	14.77	17.87			
	NT2RP3000994	13.76	10.35	18.12	10.02	5.77	7.25	4.5	8.07	7.2	*	-	
	NT2RP3001001	26.43	28.79	25.4	17.26	14.54	13.11	11.19	38.41	33.45	**	-	
	NT2RP3001004	4.86	4.98	10.62	7.73	3.07	4.03	3.13	4.4	2.41			
	NT2RP3001007	7.65	8.96	16.59	12.13	9.28	3.44	5.05	6.53	5.77			
20	NT2RP3001012	29.74	22.53	33.06	15.24	16.46	13.42	12.28	11.59	13.53	*	**	-
	NT2RP3001042	8.82	8.59	12.15	8.9	5.69	3.5	3.15	4.09	3.65	**	-	
	NT2RP3001044	9.07	11.65	10.99	8.25	9.28	5.59	7.83	7.51	5.88	*	-	
	NT2RP3001048	6.16	7.8	4.48	10.05	2.76	3.93	4.91	5.32	6.21			
	NT2RP3001050	13.37	10.61	15.08	10.7	4.83	9.6	6.34	11.26	9.4			
	NT2RP3001055	13.05	10.48	21.89	13.3	7.76	7.17	10.39	12.83	10.26			
25	NT2RP3001057	402.89	324.38	250.29	272.31	285.99	209.38	216.17	328.86	144.63			
	NT2RP3001061	7.38	5.55	10.57	7.56	3.5	2.09	3.67	3.84	2.18	*	-	
	NT2RP3001069	5.39	4.71	9.6	4.84	1.4	2.62	2.39	2.57	3.49			
	NT2RP3001074	12.1	8.82	12.25	8.96	3.61	6.95	9.25	6.39	6.94			
	NT2RP3001078	13.01	11.28	14.67	14.05	9.24	8.9	10.46	7.65	9.46	*	-	
	NT2RP3001081	8.54	6.06	5.96	10.22	3.89	7.03	6.21	3.36	4.4			
30	NT2RP3001084	37.48	37.33	40.93	15.51	10.29	21.45	23.22	30.25	29.17	**	*	-
	NT2RP3001095	7.76	24.69	13.38	9.04	3.7	5.9	3.38	7.52	4.23			
	NT2RP3001096	10.65	10.08	12.53	7.44	5.06	4.16	3.27	5.66	3.18	*	**	-
	NT2RP3001097	16.83	38.99	30.48	15.02	9.07	10.26	12.7	8.94	6.97	*	-	
	NT2RP3001107	28.61	24.48	44.48	22.05	25.16	18.26	10.04	8.83	11.02	*	-	
	NT2RP3001109	17.96	17.48	18.06	10.7	9.3	6.66	7.73	5.5	3.91	**	**	-
35	NT2RP3001111	10.67	15.58	14.57	9.33	6.41	6.38	8.41	3.97	4.53	*	*	-
	NT2RP3001112	31.69	26.22	26.11	22.8	9.31	16.42	16.88	9.36	14.31	**	-	
	NT2RP3001113	14.44	9.43	17.91	10.99	5.21	10.46	6.16	6.79	9.04			
	NT2RP3001115	254.68	276.58	258.3	135.39	153.31	123.7	118.47	299.67	339.02	**	-	
	NT2RP3001116	5.38	8	8.83	6.23	4.02	2.69	4.29	3.33	3.99	*	-	
40	NT2RP3001119	18.64	22.98	19.35	7.26	6.2	5.92	8.84	7.37	9.08	**	**	-
	NT2RP3001120	6.46	9.54	10.55	5.21	2.06	2.39	3.48	2.35	2.85	*	**	-
	NT2RP3001126	20.13	19.18	15.24	6.77	6.2	4.55	4.46	2.87	5.52	**	**	-
	NT2RP3001127	348.14	574.68	264.45	161.21	220.83	168.52	563.89	269.72	342.5			
	NT2RP3001133	93.42	110.08	112.43	155.23	89.96	123.37	108.06	86.85	85.22			
	NT2RP3001140	96.94	96.61	111.2	55.83	46.1	51.01	55.61	103.28	84.66	**	-	
	NT2RP3001147	4.72	7.64	8.71	6.37	2.09	3.65	2.74	3.66	2.93	*	-	
45	NT2RP3001150	11.47	12.3	12.64	9.98	10.24	3.21	4.96	7.94	6.36	**	-	
	NT2RP3001152	36.54	72.08	61.62	66.57	55.9	58.11	48.19	41.6	53.66			
	NT2RP3001155	19.7	23.76	17.71	11.79	7.15	8.62	14.02	11.36	10.16	**	*	-
	NT2RP3001156	19.46	20.53	21.5	20.77	20.49	12.08	20.09	17.5	18.2			
	NT2RP3001159	18.55	19.29	22.76	15.71	9.43	11.61	19.86	20.4	20.19	*	-	
	NT2RP3001170	16.8	14.86	12.69	11.85	4.5	6.68	9.82	3.85	4.53	*	*	-
50	NT2RP3001176	17.96	17.32	14.24	8.81	7.86	12.88	9.17	14.49	11.22	*	-	
	NT2RP3001195	197.73	228.27	190.42	299.15	260.22	386.11	194.62	191.1	184.16	*	+	
	NT2RP3001209	74.83	67.88	81.64	88.53	60.41	90.53	81.32	84.97	83.11			
	NT2RP3001214	17.39	13.69	27.12	17.15	13.89	13.51	7.67	14.21	13.2			
	NT2RP3001216	18.74	16.38	19.13	19.54	12.84	10.16	26.01	26.17	21.52	*	+	
	NT2RP3001221	6.31	7.39	8.65	9.62	5.18	6.37	8.56	6.05	7.92			
55	NT2RP3001226	22.75	26.78	31.16	8.49	9.07	6.76	11.49	9.21	9.9	**	**	-
	NT2RP3001230	6.51	6.56	3.99	9.46	3.93	5.31	7.3	6.36	5.9			
	NT2RP3001232	6.36	3.81	12.04	7.09	3.34	3.56	1.97	4.41	3			

Table 439

	NT2RP3000596	8.71	11.87	13.23	5.92	4.35	3.28	4.65	3.35	4	*	**	-	-
	NT2RP3000599	39.3	60.28	67.05	35.1	44.12	24.38	35.82	36.31	35.96				
5	NT2RP3000603	69.47	91.4	87.14	91.5	86.56	93.49	64.94	61.32	72.5				
	NT2RP3000605	22.45	27.45	25.73	13.14	8.89	13.46	17.63	11.14	9.76	**	*	-	-
	NT2RP3000607	10.48	11.53	16.41	10.77	7.11	9.56	5.29	9.79	8.32				
	NT2RP3000616	11.96	13.13	17.29	7.98	6.33	4.22	6.69	4.25	4.27	*	**	-	-
	NT2RP3000621	218.49	188.17	246.48	222.59	168.47	221.1	202.08	168.53	186.24				
	NT2RP3000622	11.99	8.26	9.27	9.15	3.89	5.28	6.73	6.59	3.57				
10	NT2RP3000624	27.07	18.6	22.96	16.72	12.03	13.8	17.33	23.33	18.89	*		-	
	NT2RP3000628	27.69	34.41	73.45	22.73	20.51	22.18	15.98	19.75	18.56				
	NT2RP3000631	6.03	12.91	11.07	11.79	9.91	5.14	4.52	6.12	7.95				
	NT2RP3000632	16.5	21.72	26.01	18.26	13.03	16.89	15.73	14.14	13.67				
	NT2RP3000638	31.94	42.93	38.07	22.76	17	20.23	24.56	32.44	30.42	**		-	
	NT2RP3000644	26.66	26.83	24.84	16.68	14.97	15.79	28.13	27.03	30.11	**		-	
15	NT2RP3000645	12.37	13.95	21.96	14.87	10.1	8.68	7	6.44	7.2	*		-	
	NT2RP3000652	20.17	23.26	25.45	12.67	9.39	13.26	9.11	9.19	15.37	**	*	-	-
	NT2RP3000658	6.96	4.69	8.52	6.05	2.7	4.41	2.15	3.59	3.85	*		-	
	NT2RP3000660	21.94	12.21	11.63	11.36	9.42	9.56	9.02	8.73	8.05				
	NT2RP3000661	15.39	18.23	27.58	11.62	4.78	9.25	6.26	7.03	6.07	*	*	-	-
	NT2RP3000665	34.94	43.07	59.18	27.69	16.86	21.15	16.2	20.9	14.32	*	*	-	-
20	NT2RP3000676	11.53	4.79	8.95	6.6	4.34	2.4	4.94	2.78	3.68				
	NT2RP3000677	12.19	14.6	7.18	13.97	4.51	9.55	9.82	11.58	12.06				
	NT2RP3000681	7.72	7.98	7.04	6.12	2.54	4.33	4.62	3.23	3.26	*	**	-	-
	NT2RP3000683	16.43	8.63	13.69	14.68	12.94	13.49	5.83	9.27	7.59				
	NT2RP3000685	8.41	6.46	7.19	10.24	3.04	4.6	5.55	3.8	3.45	*		-	
	NT2RP3000690	11.52	7.55	10.23	15.03	9.13	13.25	14.85	17.96	14.92	*		+	
25	NT2RP3000692	6.94	6.35	7.8	8.41	2.13	5.63	3.39	7.51	5.51				
	NT2RP3000708	9.74	10.33	12.77	11.25	8.14	10.99	9.73	10.17	12.86				
	NT2RP3000719	9.86	16.24	14.31	8.23	2.38	5.7	5.89	3.68	4.39	*	*	-	-
	NT2RP3000721	8.23	3.7	5.33	7.95	1.94	5.33	6.54	5.47	6.13				
	NT2RP3000728	4.68	4.59	4.86	7.19	1.44	3.15	5.16	3.82	3.57				
30	NT2RP3000730	14.89	15.22	17.78	16.76	11.05	11.99	16.08	10.04	9.6				
	NT2RP3000733	62.46	50.63	81.88	55.7	39.3	45.85	39.59	39.01	40.88				
	NT2RP3000735	4.91	3.22	9.89	6.52	1.99	3.91	2.03	3.65	2.09				
	NT2RP3000736	6.36	3.45	10.98	8.89	2.09	4.31	3.75	3.03	2.56				
	NT2RP3000739	40.42	44.03	40.83	24.77	27.39	34.23	25.37	20.3	32.14	*	*	-	-
	NT2RP3000742	4.84	6.47	8.58	7.03	4.15	2.9	3.28	5.74	3.55				
35	NT2RP3000753	8.3	6.8	12.66	10.6	4.35	5.47	5.99	5.75	7.45				
	NT2RP3000759	36.84	70.63	56.01	36.02	19.41	27.88	8.09	6.25	5.44	**		-	
	NT2RP3000789	45.54	27.39	34.07	33.12	27.39	34.54	20.51	27.67	27.21				
	NT2RP3000815	22.33	13.62	22.75	12.02	10.68	14.79	18.74	20.82	19.96				
	NT2RP3000818	30.57	21.11	33.63	24.24	17.69	21.68	13.73	17.32	15.5	*		-	
	NT2RP3000820	35.87	28.12	43.92	32.81	27.98	25.05	21.15	26.93	23.2				
40	NT2RP3000821	3.51	4.2	7.15	6.21	2.3	3.01	3.82	2.8	1.62				
	NT2RP3000825	16.22	23.18	33.6	11.67	17.67	13.25	9.13	8.99	16.22				
	NT2RP3000826	57.22	56.7	80.61	55.4	48.67	48.17	57.78	64.97	54.17				
	NT2RP3000836	3.1	3.53	3.85	7.53	2.33	2.42	3.69	4.81	2.68				
	NT2RP3000838	14.56	14.22	18.93	20.82	12.46	10.78	11.08	17.41	54.13				
	NT2RP3000839	45.25	36.9	44.36	35.78	27.32	20.79	19.28	33.14	25.15	*	*	-	-
45	NT2RP3000841	12.21	10.37	21.1	12.42	10.04	9.18	8.34	11.21	7.12				
	NT2RP3000845	8.59	31.26	15.5	9.69	12.93	10.25	5.37	12.21	8.92				
	NT2RP3000847	12.91	10.6	18.85	11.06	8.9	9.57	6.7	5.47	6.54	*		-	
	NT2RP3000848	11.47	16.2	11.34	8.1	7	4.83	3.86	7.4	3.83	*	*	-	-
	NT2RP3000850	21.09	31.1	29.31	19.76	12.79	16.25	10.7	10.54	5.19	*	**	-	-
	NT2RP3000852	2.73	3.9	3.17	7.51	3.17	2.39	4.15	2.97	3.66				
50	NT2RP3000859	32.3	21.43	36.02	18.9	21.77	13.31	16.61	18.23	32.52				
	NT2RP3000861	15.56	12.49	16.59	9.02	5.36	4.49	4.53	5.88	27.25	**		-	
	NT2RP3000862	12.15	10.89	17.99	10.36	6.14	6.17	5.59	8.92	5.21	*		-	
	NT2RP3000865	9.02	6.17	10.68	7.46	4.29	4.01	2.73	4.35	3.2	*		-	
	NT2RP3000866	43.46	38.1	57.47	39.52	40.18	44.87	37.48	37.57	27.56				
	NT2RP3000868	10.55	23.22	17.15	7.33	6.65	4.58	6.84	9.1	10.81	*		-	
55	NT2RP3000869	5.78	11.8	14.52	9.41	10.86	7.13	6.87	5.05	7.12				
	NT2RP3000871	184.48	180.18	141.72	252.15	170.88	170.92	247.35	193.7	214.46				
	NT2RP3000875	14.37	14.46	23.84	13.76	12.05	11.47	7.45	6.27	6.52	*		-	

Table 438

	NT2RP3000292	73.41	53.56	62.96	43.77	42.72	40.98	49.91	45.23	84.29	*	-
	NT2RP3000299	6.1	3.92	13.07	7.5	3.65	4.11	4.21	4.47	3.64		
5	NT2RP3000304	25.73	24.35	31.49	22.39	17.08	15.95	20.38	9.68	10.08	*	-
	NT2RP3000310	21.09	16.82	33.26	14.15	13.15	10.27	8.12	6.37	6.54	*	-
	NT2RP3000312	0.99	6.75	7.52	6.17	1.98	0.95	3.63	3.7	0.46		
	NT2RP3000320	11.54	11.91	24.03	15.6	10.14	7.39	6.78	13.07	6.3		
	NT2RP3000322	6.3	10.23	13.74	8.18	5.89	5.4	11.65	9.13	4.6		
	NT2RP3000324	3.35	4.35	17.76	5.33	3.9	3.91	1.03	4.83	1.94		
10	NT2RP3000326	64.19	41.44	38.83	20.52	19.15	16.6	15.62	30.96	63.08	*	-
	NT2RP3000329	11.55	11.02	22.98	13.53	7.92	12.17	10.06	11.03	12.62		
	NT2RP3000330	7.65	5.98	10.92	7.41	2.84	4.65	3.07	4.04	3.9	*	-
	NT2RP3000333	5.32	5.01	9.94	8.43	4.88	3.17	4.98	3.48	4.27		
	NT2RP3000341	13.68	14.31	17.57	13.78	9.61	9.9	9.7	9.67	11.82	*	-
	NT2RP3000344	45.81	52.79	53.7	28.28	29.87	22.15	36.08	31.27	32.79	**	**
15	NT2RP3000345	9.05	22.29	17.79	14.81	9.64	10.42	11.7	5.94	6.21		
	NT2RP3000348	152.75	76.47	143.51	181.96	103.16	145.4	96.65	108.56	99.33		
	NT2RP3000350	14.44	9.79	17.94	12.6	7.91	9.69	10.66	15.9	13.61		
	NT2RP3000359	12.38	12.66	23.75	11.88	10.77	13.58	14.05	17.04	16.51		
	NT2RP3000361	30.21	20.92	25.69	12.5	10.99	10.9	7.28	12.88	9.35	**	**
20	NT2RP3000366	69.71	53.96	58.74	44.82	38.68	40.33	27.58	41.09	30.61	*	*
	NT2RP3000378	5.8	12.66	6.1	6.74	3.98	3.58	4.72	5.63	5.68		
	NT2RP3000384	26.45	19.42	18.46	13.21	9.98	7.79	8.34	6.32	5.93	*	**
	NT2RP3000389	22.1	25.13	28.16	17.61	19.95	18.45	23.81	18.85	25.81	*	-
	NT2RP3000393	33.85	14.06	19.19	10.32	8.44	17.76	6.01	7.4	12.32		
	NT2RP3000395	24.46	28.99	21.1	13.42	10.29	14.47	9.05	14.21	24.87	**	-
	NT2RP3000397	24.89	21.35	36.91	14.31	14.82	16.14	17.25	23.74	23.2		
25	NT2RP3000398	9.55	10.75	17.31	7.25	6.82	4.6	4.16	4.17	3.99	*	-
	NT2RP3000403	93.32	64.65	56.89	57.84	43.74	45.93	35.74	40.63	30.88	*	-
	NT2RP3000418	14.37	22.01	20.2	11.88	12.29	11.51	7.67	14.89	12.11	*	-
	NT2RP3000424	40.86	56.6	51.72	39.57	32.99	30.24	31.05	29.26	23.73	*	*
	NT2RP3000427	19.59	24.95	16.59	14.3	10.74	15.64	9.16	9.9	12.52	*	-
30	NT2RP3000431	16.18	14.67	25.82	11.73	10.4	20.33	8.15	16.47	13.08		
	NT2RP3000433	25.31	54.08	36.8	35.67	31.56	27.85	24.04	39.4	34.85		
	NT2RP3000436	9.62	10.09	16.49	6.56	4.89	5.29	2.88	5.01	2.75	*	*
	NT2RP3000439	44.72	47.78	44.4	50.96	50.94	30.43	33.38	44	42.05		
	NT2RP3000441	10.82	8.95	14.82	10.83	6.19	5.02	6.65	8.63	7.39		
	NT2RP3000444	17.45	14.75	26.3	14.55	11.35	9.32	12.21	11.85	12.68		
35	NT2RP3000448	6.61	7.47	11.97	6.37	5.01	4.15	4.85	3.6	5.55		
	NT2RP3000449	17.38	26.19	23.45	8.67	7.34	7.86	3.61	4.2	2.92	**	**
	NT2RP3000451	31.29	40.71	35.46	10.79	20.02	15.29	13.25	22.78	15.67	**	**
	NT2RP3000456	6.24	4.2	13.64	8.03	3.84	0	2.84	3.63	2.55		
	NT2RP3000460	10.33	10.39	17.07	7.27	4.9	3.93	5.19	5.56	4.78	*	*
	NT2RP3000471	11.88	7.85	23.63	10.17	6.71	3.54	3.68	5.28	5.09		
40	NT2RP3000477	49.73	54.72	58.77	38.09	33.82	22.58	25.2	26.62	28.76	*	**
	NT2RP3000478	6.95	4.02	8.07	5.51	2.95	1.38	1.83	1.8	1.91	*	-
	NT2RP3000481	11.03	7.58	11.93	11.71	8.13	6.39	6.65	9.77	6.96		
	NT2RP3000484	25.46	33.21	34.57	19.56	15.47	13.67	14.97	11.89	12.9	*	**
	NT2RP3000487	21.14	29.38	33.76	13.62	12.2	14.9	5.33	12.46	12.14	*	*
	NT2RP3000512	12.44	14.88	14.98	11.55	7.87	5.16	5.05	7.04	5.14	*	**
45	NT2RP3000523	93.94	104.88	80.32	71.64	44.61	35.48	20.12	30.84	18.65	*	**
	NT2RP3000526	7.94	7.79	12.87	7.19	2.92	3.56	4	5.42	4.09	*	-
	NT2RP3000527	42.97	38.74	42.72	25	28.15	22.19	19.81	21.07	11.92	**	**
	NT2RP3000531	20.64	13.92	14.48	12.89	10.75	4.81	8.04	4.79	4.56	*	-
	NT2RP3000532	5.33	6.65	5.17	4.82	3.38	2.17	2.24	1.62	4.13	*	-
	NT2RP3000542	21.28	18.32	16.2	8.3	8.11	6.12	3.53	3.83	3.8	**	**
50	NT2RP3000554	11.71	17.67	14.03	9.45	3.02	6.99	4.57	4.9	5.09	*	**
	NT2RP3000561	21.04	16.88	23.96	14.27	9.57	9.4	4.67	15.18	7.79	*	*
	NT2RP3000562	56.42	51.6	61.28	47.36	46.18	42.62	32.74	44.25	48.68	*	-
	NT2RP3000578	19.29	21.22	25	14.6	13.72	9.4	9.07	9.7	10.06	*	**
	NT2RP3000582	9.4	7.64	13.09	8.29	5.67	6.43	5.12	6.71	5.65		
	NT2RP3000584	22.71	44.85	35.97	15.47	26.23	10.78	9.11	11.41	6.79	*	-
55	NT2RP3000586	8.23	4.88	7.14	6.37	3.31	2.97	3.91	1.9	2.63	*	-
	NT2RP3000590	7.21	5.97	5.97	5.32	2.92	1.94	4.92	2.84	1.77	*	-
	NT2RP3000592	69.63	37.92	90.02	128.75	103.66	125.15	70.02	69.72	60.26	*	+

Table 437

	NT2RP3000011	14.9	6	15.29	9.33	4.35	7.79	4.33	10.84	6.28				
	NT2RP3000014	108.09	109.54	95.92	98.76	68.18	81.91	47.71	112.2	132.8				
5	NT2RP3000016	14.35	13.63	24.71	7.65	8.46	5.45	3.86	7.49	4.11	*	*	-	-
	NT2RP3000022	7.65	10.24	13.67	9.49	4.92	2.16	2.89	2.32	2.03	*	*	-	-
	NT2RP3000024	12.64	12.37	14.37	11.32	5.72	6.81	6.24	6.05	6.52	*	**	-	-
	NT2RP3000031	77.24	79.76	145.67	144.42	128.63	86.96	77.23	69.72	83.66				
	NT2RP3000034	6.27	6.62	8.74	6.38	3.4	3.43	4.64	3.79	2.96	*	*	-	-
	NT2RP3000037	17.58	19.44	20.81	11.72	11.55	13.9	18.45	10.67	14.15	**	*	-	-
10	NT2RP3000040	27.59	16.23	24.01	13.53	7.79	10.55	2.51	8.1	6.81	*	*	-	-
	NT2RP3000041	5.08	4.46	12.76	6.9	3.15	3.65	6.44	3	1.87				
	NT2RP3000046	7.63	5.75	15.27	8.29	3.02	2.83	2.71	4.54	5.05				
	NT2RP3000047	5.93	4.81	14.06	5.49	2.34	2.87	0.9	3.18	1.51				
	NT2RP3000049	6.45	8.2	10.35	6.13	1.6	1.78	2.62	3.99	1.9	*	*	-	-
	NT2RP3000050	11.72	13.02	12.95	7.97	7.04	4.74	3.27	4.45	3.93	**	**	-	-
15	NT2RP3000051	20.55	9.33	18.87	14.09	17.37	4.84	3.99	4.5	4.21	*	*	-	-
	NT2RP3000054	6.46	5.61	6.8	6.67	3.52	2.38	1.52	2.23	2.77	**	*	-	-
	NT2RP3000055	18.47	13.61	17.33	12.33	8.89	12.37	7.71	11.31	9.35	*	*	-	-
	NT2RP3000056	1.67	2.36	8.51	3.35	1.42	1.09	0.81	1.36	0.36				
	NT2RP3000059	7.89	7.9	22.29	10.9	7.16	5.13	2.59	4.88	2.31				
	NT2RP3000063	1.84	4.14	7.63	4.56	1.09	0.43	0.47	0.97	0				
20	NT2RP3000068	6.17	8.89	16.21	5.26	4.81	2.29	6.13	4.42	2.89				
	NT2RP3000069	9.15	8.31	8.75	6.58	4.56	4.92	4.17	3.76	6.03	**	**	-	-
	NT2RP3000072	32.05	40.26	39.15	31.11	26.2	25.34	26.47	26.26	26.37	*	*	-	-
	NT2RP3000080	25.2	23.2	22.56	18.48	15.53	13.5	21.79	14.38	13.52	**	*	-	-
	NT2RP3000085	67.96	43.89	70.79	53.46	33.71	65.94	52.26	69.62	62.27				
25	NT2RP3000087	111.9	96.79	117.72	81.88	54.53	62.52	49.82	71.54	66.46	*	**	-	-
	NT2RP3000092	9.87	12.43	24.83	8.97	4.39	4.92	2.95	7.36	3.67				
	NT2RP3000109	10.57	9.88	12.54	8.31	8.64	5.99	6.65	6.37	5.22	*	**	-	-
	NT2RP3000119	8.08	6.85	10.03	6.29	4.73	2.45	4.69	3.89	2.89	*	*	-	-
	NT2RP3000125	6.75	8.13	6.71	5.47	2.88	2.09	0.61	2.51	2.19	*	**	-	-
	NT2RP3000131	13.6	16.29	13.49	12.52	5.73	6.16	5.37	4.48	4.22	**	**	-	-
30	NT2RP3000134	18.43	21.97	18.23	9.28	9.9	8.55	6.96	2.89	3.6	**	**	-	-
	NT2RP3000137	10.23	18.11	17.39	15.98	4.55	11.71	4.48	6.98	5.44	*	*	-	-
	NT2RP3000142	35.53	27.58	35.61	21.77	15.45	19.48	21.05	30.25	26.9	*	*	-	-
	NT2RP3000148	26.32	39.38	34.66	19.53	19	15.35	18.54	17.46	14.71	*	*	-	-
	NT2RP3000149	32.9	38.83	27.1	14.1	20.26	8.54	10.04	7.73	9.34	*	**	-	-
	NT2RP3000163	8.85	9.04	13.21	8.67	3.56	5.95	3.7	3.77	3.35	**	*	-	-
35	NT2RP3000168	15.39	20.39	19.23	9.48	11.8	5.55	7.63	15.26	9.88	*	*	-	-
	NT2RP3000169	15.25	12.33	13.06	7.58	7.27	4.09	8.07	3.27	4.36	**	**	-	-
	NT2RP3000171	6.75	12.8	10.62	11.26	6.03	4.95	8.98	2.23	5.04				
	NT2RP3000172	40.04	25.12	32.12	25.17	9.97	24.6	26.83	33.18	37.52				
	NT2RP3000186	16.1	16.13	19.67	12.39	6.63	8.09	5.65	10.05	10.55	*	*	-	-
	NT2RP3000197	50.94	62.27	70	34.47	33.63	20.62	34.62	48.18	55.04	*	*	-	-
40	NT2RP3000201	14.53	11.88	22.5	7.11	4.59	3.11	5.14	5.79	5.41	*	*	-	-
	NT2RP3000204	4.16	4.96	6.59	4.48	1.18	2.3	2.66	1.7	2.27	*	*	-	-
	NT2RP3000207	95.59	249.91	234.63	30.21	59.19	29.76	35.62	12.46	31.91	*	*	-	-
	NT2RP3000216	7.95	7.31	6.95	6.17	2.69	2.5	2.94	3.29	3.98	*	**	-	-
	NT2RP3000220	22.18	29.86	28.92	10.43	5.88	15.01	13.77	5.33	14.12	**	*	-	-
	NT2RP3000221	10.31	9.17	11.73	12.25	5.03	10.36	5.52	10.77	5.91				
45	NT2RP3000232	46.66	42.43	53.26	15.69	8.2	7.9	5.53	11.37	6.25	**	**	-	-
	NT2RP3000233	8.3	12.35	2.97	7.16	16.65	2.19	9.24	3.42	2.61				
	NT2RP3000234	3.83	3.88	5.86	5.35	3.32	2.04	1.6	2.52	2.2	*	*	-	-
	NT2RP3000235	9.35	15.39	13.27	13.07	5.58	8.43	10.72	9.32	10.97				
	NT2RP3000239	18.63	25.23	22.23	15.58	11.06	7.77	9.06	9.28	12.71	*	**	-	-
	NT2RP3000247	32.06	44.16	41.62	22.69	17.24	20.67	24.3	20.29	19.85	**	*	-	-
50	NT2RP3000251	44.56	41.2	38.69	25.94	14.62	23.41	43.42	39.09	44.51	**	*	-	-
	NT2RP3000252	21.9	16.55	21.84	18.88	16.64	19.17	13.44	23.92	18.04				
	NT2RP3000255	18.67	16.02	17.43	18.73	10.56	14.35	9.36	10.73	8.47	**	*	-	-
	NT2RP3000262	17.74	15.84	18.49	25.1	16.29	24.75	12.36	12.76	13.36	**	*	-	-
	NT2RP3000266	37.8	22.61	54.63	46.91	29.06	37.64	30.75	39.28	44.8				
	NT2RP3000267	5.43	5.61	4.39	6.25	5.05	2.74	3.18	2.68	2.59	**	*	-	-
55	NT2RP3000271	4.92	10.2	10.62	7.08	4.11	3.49	3.88	5.36	2.72				
	NT2RP3000276	8.77	11.93	8.85	9.67	5.55	4.79	8.29	7.56	5.48				
	NT2RP3000281	32.19	29.94	32.48	16.28	8.76	14.6	16.94	10.41	13.64	**	**	-	-

Table 436

	NT2RP2006134	9.57	6.53	12.45	11.76	2.26	6.54	9.28	6.55	5.96			
	NT2RP2006141	13.6	10.51	20.75	11.91	8.44	5.24	5.17	8.6	15.73			
5	NT2RP2006166	43.39	37.58	47.62	23.75	15.94	0.88	44.65	56.53	70.7	*		
	NT2RP2006176	17.92	13.68	20.98	13.87	7.31	8.84	7.42	10.41	8.82	*	*	
	NT2RP2006181	4.75	8.83	6.23	3.92	1.81	1.35	3.97	2.81	2.8	*		
	NT2RP2006184	64.52	54.32	63.63	42.14	32.28	42.22	63.72	47.39	43.75	**		
	NT2RP2006186	8.19	11.92	9.4	6.19	2.49	2.83	2.18	2.13	1.87	*	**	
	NT2RP2006196	22.53	23.16	21.9	14.52	7.82	10.37	16.26	11.79	10.32	**	**	
10	NT2RP2006199	9.01	9.11	8.26	8.72	4.43	4.65	6.19	4.21	5.05	**		
	NT2RP2006200	7.28	4.32	9.53	5.05	3.47	3.37	3.59	4.46	2.65			
	NT2RP2006210	90.78	69.8	53.22	55.44	37.83	25.49	20.22	40.51	30.76	*		
	NT2RP2006219	11.28	20.39	17.51	8.1	14.35	6.63	11.34	9.44	19.6			
	NT2RP2006224	12.75	19.13	21.38	12.72	10.81	3.64	17.87	16.61	1.85			
	NT2RP2006237	7.47	20.77	9.96	5.08	2.48	0.88	3	2.56	0.6			
15	NT2RP2006238	10.52	14.31	12.03	6.19	1.38	2.03	2.55	2.62	2.66	**	**	
	NT2RP2006258	10.05	20.38	11.9	6.71	3.4	2.58	5.8	4.54	6.64	*		
	NT2RP2006261	8.38	9.18	5.31	9.6	2.51	4.78	2.42	4.07	4.58	*	*	
	NT2RP2006269	17.03	14.29	15.32	11	5.59	7.26	3.86	12.06	5.85	*	*	
	NT2RP2006275	46.12	39	46.05	70.83	38.96	80.2	37.5	48.02	38.3			
	NT2RP2006282	18.4	26.43	24.55	13.47	12.52	6.62	5.66	6.21	6.64	*	**	
20	NT2RP2006302	17.18	26.6	32.5	13.26	6.56	8.01	5.28	4.17	5.65	*	*	
	NT2RP2006312	31.59	23.04	41.51	9.42	5.96	4.21	6.1	4.79	7.53	*	**	
	NT2RP2006320	41.28	41.71	36.93	24.86	13.12	15.23	28.98	37.1	32.63	**		
	NT2RP2006321	10.41	9.33	12.86	10.26	2.43	5.6	7.6	9.6	8.78			
	NT2RP2006323	7.58	5.4	4.29	6.62	1.35	3.01	7.41	4.95	4.91			
	NT2RP2006333	5.03	7.52	4.99	6.18	1.44	5.34	2.1	4.81	3.82			
25	NT2RP2006334	7.94	8.71	7.64	8.5	4.46	6.47	2.3	4.34	3.32	**		
	NT2RP2006338	12.7	9.34	36.56	7.32	2.93	4.16	2.16	2.04	3.64			
	NT2RP2006339	6.41	4.31	4.61	6.37	3.36	3.67	2.31	2.9	2.66	*		
	NT2RP2006355	6.78	12.38	3.94	6.81	1.66	2.59	2.74	4.14	2.84			
	NT2RP2006365	5.27	2.38	4.32	5.39	0.85	1.53	2.13	3.92	2.48			
30	NT2RP2006374	123.01	124.5	151.43	183.62	144.52	148.4	137.08	112.64	166.86			
	NT2RP2006393	22.69	18.53	23.1	17.79	6.84	13.61	27.67	23.54	25.74			
	NT2RP2006394	15.6	11.55	17.73	16.97	12.66	13.34	9.59	13.69	11.86			
	NT2RP2006400	8.82	4.99	12	10.48	4.15	11.24	9.88	11.64	10.5			
	NT2RP2006411	4.42	4.9	11.08	5.28	2.3	3.07	2.25	3.04	3.21			
	NT2RP2006429	2.69	1.8	8.38	6.1	1.17	1.68	2.03	2.68	1.46			
35	NT2RP2006435	1.14	1.44	3.96	3.86	0	1.28	1.25	0.25	0.25			
	NT2RP2006436	18.89	25.65	29.36	23.03	10	15.2	18.88	16.71	17.66			
	NT2RP2006441	3.24	5.37	8.45	6.78	2.17	3.47	5.5	2.69	1.87			
	NT2RP2006447	19.2	14.3	18.35	21.17	15.45	16.45	6.09	13.77	11.65			
	NT2RP2006454	8.01	2.56	13.95	6.26	3.2	3.65	3.95	4.09	2.96			
	NT2RP2006455	37.87	46.16	39.71	20.34	20.45	22.87	31.23	29.73	37.11	**		
40	NT2RP2006456	12.66	10.34	20.88	11.22	5.29	5.49	7.54	4.09	3.92			
	NT2RP2006464	49.31	38.6	70.84	30.42	38.8	32.65	22.89	19.33	15.74	*		
	NT2RP2006467	7.41	7.72	13.37	8.13	6.48	4.56	10.4	4.81	8.63			
	NT2RP2006472	17.27	35.33	39.96	36.59	25.1	29.12	15.84	31.69	29.65			
	NT2RP2006474	5.31	10.55	8.31	9.45	5.79	7.02	7.91	11.16	8.64			
	NT2RP2006475	14.67	7.33	18.42	13.4	5.52	5.53	5.81	10.03	7.38			
45	NT2RP2006476	21.17	14.25	23.98	17.93	10.29	15.83	16.59	21.36	17.75			
	NT2RP2006501	80.64	47.79	67.54	47.86	36.74	36.28	29.87	41.76	52.91			
	NT2RP2006512	10.73	9.4	12.56	6.68	7.05	5.5	3.87	4.91	5.32	*	**	
	NT2RP2006526	12.09	17.75	22.2	19.62	16.43	13.33	17.31	20.34	13.23			
	NT2RP2006527	4.41	6.17	7.16	7.48	1.84	1.55	3.51	2.56	2.84	*		
	NT2RP2006534	4.37	9.41	7.99	5.05	4.19	2.96	4.68	2.83	2.5			
50	NT2RP2006537	5.9	9.98	11.68	11.28	5.12	7.83	10.34	5.58	8.5			
	NT2RP2006543	12.11	8.37	13.04	12.23	3.26	11.04	6.8	5.53	9.23			
	NT2RP2006554	7.01	5.37	13.82	7.47	2.64	2.23	3.56	4.89	9.98			
	NT2RP2006565	14.34	12.69	22.07	6.94	4.33	4.74	3.36	7.61	8.3	*	*	
	NT2RP2006571	65.34	49.49	69.38	45.6	41.55	34.63	47.98	65.86	43.88	*		
	NT2RP2006573	12.86	9.58	16.76	8.82	7.97	5.87	5.7	4.86	5.46	*		
55	NT2RP2006588	5.92	15.24	10.14	5.39	4.48	3.2	2.52	3.49	4.84			
	NT2RP2006601	5.92	8.92	7.57	6.45	4.69	2.53	4.15	3.79	3.98	*		
	NT2RP3000002	17.28	31.41	14.62	22.94	26.64	18.92	27.08	23.91	17.49			

Table 435

	NT2RP2005737	61.08	46.51	54.73	57.66	30.52	54.9	80.55	70.19	69.8	*	+
	NT2RP2005741	8.17	4.22	11.68	8.13	3	5.42	3.62	4.85	3.95		
	NT2RP2005748	8.36	5.27	13.52	8.19	3.71	6.33	3.46	4.51	7.44		
5	NT2RP2005752	31.75	22.62	38.43	14.99	13.75	18.04	15.51	16.74	18.45	*	*
	NT2RP2005753	34.44	30.95	48.77	33.24	26.02	18.11	26.19	25.32	15.55		
	NT2RP2005763	8.78	7.62	12.88	6.76	3.85	3.68	4.32	5.6	4.17	*	
	NT2RP2005767	8.89	8.69	15.86	7.42	5.16	5.67	6.13	4.89	5.16		
	NT2RP2005773	53.73	61.68	69.19	40.66	32.98	33.21	37.38	30.92	21.66	**	**
10	NT2RP2005774	13.75	12.5	18.48	16.31	7.06	7.63	18.33	17.56	17.06		
	NT2RP2005775	12.13	7.53	22.24	13.73	4.82	6.66	4.04	6.6	5.74		
	NT2RP2005781	39.68	23.61	38.05	24.73	22.45	22.65	9.91	16.91	23.06		
	NT2RP2005784	30.34	20	41.77	17.54	13.38	14.34	11.13	13.26	15.12		
	NT2RP2005789	16.92	16.55	24.71	17.16	13.64	7.39	8.95	14.02	9.78		
	NT2RP2005799	11.23	8.75	9.48	10.01	3.28	4.24	3.37	6.03	3.88	**	*
15	NT2RP2005804	106.7	97.93	105.55	76.62	70.98	67.98	110.4	84.7	108.91	**	*
	NT2RP2005812	6.56	9.52	8.36	6.62	4.2	4.89	5.32	2.9	2.58	*	
	NT2RP2005815	2.76	4.98	5.71	6.34	2.94	2.66	4.39	4.52	4.97		
	NT2RP2005835	26.87	25.04	37.02	18.31	16.79	16.79	10.34	14.43	39.55	*	*
	NT2RP2005841	22.25	27.56	27.52	20.49	14.43	16.92	11.4	11.65	19.51	*	*
20	NT2RP2005853	18.13	15.89	21.67	12.41	10.18	1.04	9.88	13.51	11.3	*	*
	NT2RP2005857	11.03	8.75	10.83	7.42	4.51	1.32	4.21	5.92	5.5	*	**
	NT2RP2005859	12.15	11.97	15.23	11.24	7.93	4.86	8.29	8.94	6.43	*	
	NT2RP2005860	0.96	5.77	2.68	4.74	2.31	1.31	1.34	1.77	1.13		
	NT2RP2005863	8.01	8.99	7.06	7.45	6.66	5.32	7.67	5.36	4.12		
	NT2RP2005868	5.66	10.09	8.76	11.28	4.39	6.75	7.63	5.42	5.14		
25	NT2RP2005876	29.94	9.36	44.21	25.04	19.27	15.81	14.52	17.33	37.26		
	NT2RP2005878	17.4	21.07	28.07	15.86	10.5	10.77	9.96	17.05	21.58		
	NT2RP2005883	9.58	9.62	13.3	8.93	6.54	7.61	3.43	11.45	8.28		
	NT2RP2005886	109.48	86.27	131.02	108.7	95.74	145.77	77.39	72	86.69	*	**
	NT2RP2005887	12.65	18.34	15.15	9.59	5.47	2.19	3.8	5.68	5.07	*	**
	NT2RP2005890	11.6	13.15	14.84	6.76	5.32	5.53	4.7	5.77	4.32	**	**
30	NT2RP2005901	11.96	15.27	14.95	10.18	6.55	8.12	6.42	5	5.8	*	**
	NT2RP2005902	4.82	5.49	5.62	9.24	3.69	5.72	4.9	8.33	5.18		
	NT2RP2005908	24.41	23.56	59.06	23.37	17.17	15.35	18.41	25.09	27.9		
	NT2RP2005927	7.88	6.48	15.44	8.31	4.33	4.46	3.45	5.76	4.29		
	NT2RP2005933	14.92	15.67	28.13	12.37	6.74	6	4	6.32	4.28	*	
	NT2RP2005941	10.69	8.58	13.22	7.84	3.9	3.96	2.9	4.82	2.65	*	**
	NT2RP2005942	10.35	10.52	11.7	8.52	4	3.46	3.66	4.6	2.9	*	**
35	NT2RP2005946	8.12	19.6	14.91	5.77	6.58	2.97	4.94	4.86	6.09		
	NT2RP2005970	73.5	72.19	81.01	50.77	49.15	40.53	67.44	62.72	62.64	**	*
	NT2RP2005980	7.01	5.07	5.9	6.68	2.11	2.14	3.31	3.86	4.51	*	*
	NT2RP2005994	12.99	9.14	13.53	7.59	4.09	0.3	2.32	3.71	4.52	*	**
	NT2RP2006004	7.54	4.47	13.06	6.53	2.59	2.1	2.72	2.9	3.13		
40	NT2RP2006013	11.06	6.58	20.3	11.12	5.96	2.47	2.76	11.94	3.04		
	NT2RP2006023	79.4	60.83	74.67	51.53	34.64	38.07	64.39	76.12	79.93	*	
	NT2RP2006028	23.13	26.87	22.92	16.89	12.58	6.73	6.52	8.45	15.41	*	**
	NT2RP2006038	4.35	4.75	6.83	4.07	1.31	0.83	1.09	0.16	1.4	**	
	NT2RP2006042	15.45	13.74	13.53	9.24	8.86	3.55	5.34	6.22	7.22	*	**
	NT2RP2006043	17.81	13.76	15.29	13.7	7.47	8.55	9.41	5.9	6.89	**	
45	NT2RP2006052	12.36	8.55	16.5	5.69	3	4.31	2.78	3.7	3.31	*	*
	NT2RP2006057	7	6.58	13.05	6.99	3.77	3.67	1.75	3.57	2.95	*	*
	NT2RP2006064	31.45	32.66	33.8	25.4	15.66	8.71	6.36	7.35	7.36	*	**
	NT2RP2006068	7.85	6.5	16.27	7.3	3.64	3.64	2.88	4.02	3.43		
	NT2RP2006069	10.49	8.04	9.64	6.75	2.89	1.65	2.28	1.16	1.94	*	**
	NT2RP2006071	12.09	22.31	12.61	10.26	7.27	0.85	3.44	3.17	3.04	*	
50	NT2RP2006090	10.61	10.59	22.59	7.62	4.75	0.57	5.24	4.28	5.3		
	NT2RP2006092	15.25	7.07	12.08	13.1	8.69	10.62	13.15	8.79	8.66		
	NT2RP2006097	52.21	101.2	64.35	48.63	48.31	38.97	25.59	53.43	52.7		
	NT2RP2006098	13.67	12.41	21.3	14.85	6.15	8.49	5.13	8.68	3.6	*	
	NT2RP2006099	65.92	60.33	80.23	76.5	49.76	0.24	73.49	73.17	78.2		
	NT2RP2006100	14.09	12.17	17.63	9.6	8.02	0.77	4.01	5.07	4.89	**	
55	NT2RP2006103	6.26	4.22	7.72	5.83	3.58	2.06	2.74	2.72	2.55	*	
	NT2RP2006106	48	36	55.68	34.38	35.82	35.56	24.01	21.59	18.79	*	
	NT2RP2006127	6.5	4.86	5.96	5.76	1.93	3.45	3.78	1.37	2.06	*	

Table 434

	NT2RP2005429	20.77	21.25	24.75	28.87	19.9	22.9	31.5	41.11	27.5				
	NT2RP2005436	21.72	21.01	36.27	15.82	10.49	11.17	6.19	10.76	15.31				
5	NT2RP2005441	12.34	10.42	16.43	6.82	3.95	2.88	3.88	5.58	7.88	*	*	-	-
	NT2RP2005442	100.99	76.34	97.05	85.22	107.34	64.16	56.06	96.44	76.81				
	NT2RP2005444	44.56	42.84	50.52	45	41.48	37.39	33.32	34.33	40.3		*	-	-
	NT2RP2005453	6.3	5.27	7.3	6.26	2.81	2.81	2.27	2.79	2.65	**		-	-
	NT2RP2005457	65.08	106.4	103.3	51.25	79.76	59.93	40.37	56.9	47.82		*	-	-
	NT2RP2005458	19.63	18.76	13.48	12.24	12.61	6.33	8.23	12.39	11.01		*	-	-
10	NT2RP2005463	26.71	23.66	25.36	21.35	13.39	12.75	11.08	11.67	8.93	*	**	-	-
	NT2RP2005464	28.8	25.9	35.07	16.26	12.23	8.38	4.05	11.36	13.81	**	**	-	-
	NT2RP2005465	8.44	13.41	18.06	10.55	9.11	6.34	6.69	6.77	5.88				
	NT2RP2005472	156.47	127.25	179.85	146.2	122.26	139.88	141.98	194.05	159				
	NT2RP2005476	18.49	17.23	18.61	16.08	8.51	7.62	10.76	9.58	6.82	**		-	-
	NT2RP2005490	38.19	48.01	41.98	41.4	60.4	46.27	36.8	25.85	27.98	*		-	-
15	NT2RP2005491	79.97	62.21	87.46	43.7	46.69	33.15	42.26	41.55	39.38	*	**	-	-
	NT2RP2005495	8.67	12.67	12.73	8.05	5.23	3.12	3.29	2.22	4.63	*	**	-	-
	NT2RP2005496	40.82	33.68	40.13	35.53	16.36	22.99	49.27	41.88	52.93				
	NT2RP2005498	14.64	10.07	20.28	8.15	3.71	0.94	3.15	5.06	7.24	*	*	-	-
	NT2RP2005501	15.01	11.26	15.06	10.4	6.02	5.78	5.37	7.88	6.8	*	**	-	-
	NT2RP2005506	386.46	291.76	402.21	469.63	426.17	208.66	248.04	333.44	241.6				
20	NT2RP2005509	12.93	15.67	24.29	8.78	9.8	9.01	11.69	14.68	14.73				
	NT2RP2005514	15.12	18.77	16.43	9.09	7.1	7.44	5.07	6.13	3.95	**	**	-	-
	NT2RP2005520	125.71	88	119.54	107.31	84.59	104.73	102.45	122.23	104.75				
	NT2RP2005525	11.61	19.18	16.45	11.91	8.22	5.77	3.77	4.63	4.84	**		-	-
	NT2RP2005531	6.49	2.54	4.37	5.65	2.65	0.74	1.72	1.38	1.4				
25	NT2RP2005535	28.78	24.33	27.3	18.42	7.97	16.95	13.78	20.87	16.38	*	*	-	-
	NT2RP2005539	7.33	11.94	18.75	8.51	6.39	4.26	3.51	3.63	4.88				
	NT2RP2005540	16.33	11.75	18.94	9.77	17.11	5.41	9.79	8.14	4.41	*		-	-
	NT2RP2005541	14.69	11.15	19.91	13.02	7.16	8.12	7.95	8.57	7	*		-	-
	NT2RP2005549	11.12	11.63	18.04	9.08	7.83	5.49	4.23	4.44	5.21	*		-	-
	NT2RP2005555	16.74	21.24	18.63	10.59	10.43	7.67	9.05	5.08	3.67	**	**	-	-
30	NT2RP2005557	15.1	14.02	12.36	10.06	7.31	4.94	4.63	4.36	4.73	*	**	-	-
	NT2RP2005581	28.08	26.29	29.21	26.59	10.92	19.41	31.86	21.24	23.29				
	NT2RP2005586	16.13	21.05	17.82	8.84	4.94	6.39	4.6	7.07	6.9	**	**	-	-
	NT2RP2005597	6.89	6.97	9.77	7.01	1.89	3.66	2.71	4.02	1.82	*		-	-
	NT2RP2005600	10.61	7.34	12.5	8.41	8.29	5.16	5.14	5.97	3.47	*		-	-
	NT2RP2005605	53.68	52.91	55.5	30.27	27.97	29.06	25.09	21.21	26.25	**	**	-	-
35	NT2RP2005614	9.49	10.07	16.9	6.5	3.69	4.19	2.85	3.64	2.63	*	*	-	-
	NT2RP2005620	17.66	10.99	8.05	7.1	2.87	3.18	7.89	3.09	3.37				
	NT2RP2005622	16.16	16.55	15.71	9.98	5.85	5.98	5.01	2.43	3.54	**	**	-	-
	NT2RP2005632	24.79	21.59	16.99	24.01	11.37	16.72	22.78	10.85	12.69				
	NT2RP2005635	41.82	31.3	47.35	47.22	30.84	66.01	30.58	31.82	30.82				
	NT2RP2005637	8.32	23.19	12.21	8.57	5.9	4.47	3.6	7.75	5.54				
40	NT2RP2005640	15.46	12.6	16.31	14.33	9.36	13.4	14.45	17.32	12.67				
	NT2RP2005645	23.93	18.08	21.54	10.6	8.02	9.84	11.24	9.09	9.16	**	**	-	-
	NT2RP2005651	27.47	30.93	34.21	15.98	10.8	18.12	23.48	20.74	24.88	**	*	-	-
	NT2RP2005654	13.13	10.24	14.43	6.36	4.26	2.41	3.56	3.27	5.28	**	**	-	-
	NT2RP2005666	12.61	72.48	13.27	7.56	5.9	4.42	7.43	6.32	9.59				
	NT2RP2005669	15.23	15	15.44	12.86	8.17	11.55	8.81	7.49	8.17	*	**	-	-
45	NT2RP2005670	7.59	11.17	16.69	7.09	6.37	6.24	3.98	5.41	8.38				
	NT2RP2005671	8.66	9.38	12.5	6.17	4.27	6.41	3.39	7.42	4.35	*	*	-	-
	NT2RP2005675	53.41	58.69	56.21	73.47	47.08	88.08	46.13	47.46	45.27	**		-	-
	NT2RP2005683	12.5	14.86	22.54	12.03	8.9	11.13	12.51	11.13	11				
	NT2RP2005680	7.16	9.78	9.15	7.31	2.58	3.76	6.01	4.75	5.81	*		-	-
	NT2RP2005694	13.45	13.78	12.27	8.68	4.9	4.46	6.63	6.81	8.01	**	**	-	-
50	NT2RP2005701	21.29	27.54	24.8	13.54	7.14	7.23	9.96	8.67	7.38	**	**	-	-
	NT2RP2005712	5.67	6.5	6.73	7.41	2.31	3.04	4.59	2.86	2.66	*		-	-
	NT2RP2005719	17.92	12.64	12.02	8.92	5.92	13.08	4.64	11.29	9.06				
	NT2RP2005722	24.17	15.94	25.68	22.5	8.47	9.89	8.38	12.95	9.48	*		-	-
	NT2RP2005723	11.76	19.91	11.37	8.9	4.43	7.12	5.8	6.31	6.59	*		-	-
	NT2RP2005726	13.16	8.51	18.51	13.17	7.93	10.58	7.85	8.96	10.35				
55	NT2RP2005729	20.35	17.21	20.34	11.64	9.52	5.78	12.25	8.44	10.44	**	**	-	-
	NT2RP2005731	4.25	2.53	4.11	5.17	0	0.69	1.06	1.65	0.86	*	*	-	-
	NT2RP2005732	54.72	53.14	65.72	91.69	67.12	83.97	67.17	50.47	67.53	*	+		

Table 433

	NT2RP2004967	18.8	14.08	15.99	13.12	6.04	9.98	11.43	12.91	9.85	*	*	-
	NT2RP2004974	8.02	8.78	7.74	5.74	3.27	3.73	5.44	2.35	2.1	**	*	-
5	NT2RP2004978	14.12	27.31	13.7	11.96	4.65	6.79	4.16	9.57	4.35			
	NT2RP2004982	4.88	6.54	10.22	4.28	1.48	0.85	1.99	1	1.83	*		-
	NT2RP2004985	209.85	195.75	215.75	166.58	153.94	208.41	198.37	224.84	192.82			
	NT2RP2004999	16.04	12.87	26.42	17.44	8.41	10.32	9.65	18.31	9.11			
	NT2RP2005000	9.18	10.25	11.07	6.99	2.87	3.26	5.46	4.95	3.42	*	**	-
	NT2RP2005001	18.08	20.79	40.06	15.62	11.69	13.08	12.25	15.19	13.27			
10	NT2RP2005003	15.88	14.42	15.05	11.32	5.53	6.52	11.83	11.19	8.75	*	*	-
	NT2RP2005012	40.61	57.66	43.61	20.05	11.76	16.65	21.02	13.16	20.53	**	**	-
	NT2RP2005018	6.29	10.32	12.32	7.2	3.34	4.91	3.54	5.22	4.18	*		-
	NT2RP2005020	138.98	109.02	105.29	97.72	59.61	79.54	117.66	175.16	138.64			
	NT2RP2005022	8.77	8.01	12.34	8.19	7.74	5.07	4.96	7.16	9.74			
	NT2RP2005027	53.46	62.9	53.33	58.43	50.24	56.5	33.74	38.11	31.82	**		-
15	NT2RP2005031	6.85	8.81	9.58	5.73	4.02	3.36	3.46	3.48	3.14	*	**	-
	NT2RP2005037	16.82	18.46	19.47	6.47	4.36	2.93	2.65	3.66	6.14	**	**	-
	NT2RP2005038	6.96	8.98	11.09	7.2	2.46	3.49	4.34	2.45	1.1	*		-
	NT2RP2005048	39.21	28.46	29.89	31.24	22.21	43.14	32.34	46.11	37.29			
	NT2RP2005089	79.3	47.14	64.37	36.64	18.08	20.34	23.96	33.08	29.72	*	*	-
20	NT2RP2005073	67.78	54.66	62.54	70.16	48.08	81.7	55.95	55.76	57.18			
	NT2RP2005097	10.25	7.51	14.59	6.49	1.21	4.7	3.75	2.8	2.35	*		-
	NT2RP2005108	13.57	11.63	13.11	6.87	5.69	6.68	5.31	6.56	5.73	**	**	-
	NT2RP2005116	11.75	10.87	12.14	10.24	4.17	4.29	7.43	5.88	6.2	**		-
	NT2RP2005126	21.43	28.84	28.25	14.77	6.89	5.64	7.77	5.68	5.93	*	**	-
	NT2RP2005135	10.21	11.92	11.29	12.06	3.02	7.4	9.29	7.72	6.92	*		-
25	NT2RP2005139	4.47	5.9	6.41	5.94	3.02	9.46	4.16	4.52	4.09			
	NT2RP2005140	10.93	7.52	7.44	7.58	3.55	5.44	3	4.22	2.26	*		-
	NT2RP2005144	13.73	10.54	13.89	8.29	4.49	7.83	1.84	3.98	2.4	*	**	-
	NT2RP2005147	22.69	15.06	21.74	11.07	5.61	11.36	5.53	13.64	8.82	*	*	-
	NT2RP2005148	19.34	14.88	12.3	16.32	7.5	14.47	18.68	18.99	24			
	NT2RP2005159	9.52	7.06	9.83	13.34	5.14	7.92	4.36	3.56	6.05	*		-
30	NT2RP2005162	14.54	12.5	16.31	14.53	7.94	8	8.72	9.34	7.02	**		-
	NT2RP2005163	25.5	37.52	33.86	15.82	13.92	18.66	24.95	23.28	21.32	*		-
	NT2RP2005168	10.58	6.62	14.16	11.05	5.89	4.43	4.83	6.74	4.36			
	NT2RP2005181	4.05	2.99	8.75	6.74	2.11	1.29	1.74	3.19	2.03			
	NT2RP2005204	14.87	10.24	20.73	15.12	7.06	10.51	11.01	6.88	10.24			
	NT2RP2005219	24.24	19.15	32.19	26.01	12.29	15.28	21.03	18.69	23.06			
35	NT2RP2005227	12.24	9.75	17.91	17.67	5.27	7.45	11.51	12.06	11.25			
	NT2RP2005237	103.35	89.54	103.29	128.64	70.44	96.83	81.16	75.84	83.42	*		-
	NT2RP2005239	4.86	6.22	9.52	8.57	2.23	3.55	3.5	4.01	4.35			
	NT2RP2005247	22.45	26.01	32.27	32.43	14.24	19.04	34.79	31.96	33.52			
	NT2RP2005254	36.5	15.86	25.1	17.04	11.36	15.24	5.51	13.65	13.69			
	NT2RP2005270	25.5	13.21	29.5	13.51	9.93	13.53	11.24	14.21	22.67			
40	NT2RP2005276	40.6	21.43	40.17	20.18	14.43	16.86	10.58	8.43	8.36	*		-
	NT2RP2005287	16.9	10.22	25.14	13.6	9.12	7.29	8.33	9.76	8.8			
	NT2RP2005288	6.35	8.09	8.85	7.18	3.2	3.43	4.54	4.39	3.3	*		-
	NT2RP2005289	19.76	15.26	21.43	14.2	10.57	10.92	15.53	14.93	15.73	*		-
	NT2RP2005293	9.79	11.07	10.95	8.57	5.47	3.14	6.98	5.79	5.74	*	**	-
	NT2RP2005315	15.84	21	21.65	20.97	8.72	12.4	13.48	9.21	11.44	*		-
45	NT2RP2005322	33.55	31.7	65.28	25.82	25.53	26.43	14.38	22.11	31.46			
	NT2RP2005325	54.2	44	55.99	38.68	30.82	29.07	18.94	27.07	30.65	*	**	-
	NT2RP2005336	20.36	12.98	27.96	11.35	7.87	9.01	10.99	13.9	14.17			
	NT2RP2005343	12.61	22.98	14.15	9.9	5.77	8.7	7.81	10.19	10.12			
	NT2RP2005344	6.04	8.8	10.41	6.82	3.18	3.71	4.76	3.86	3.88	*		-
	NT2RP2005347	7.59	13.3	8.99	7.33	4.35	4.03	4.9	3.27	3.77	*		-
50	NT2RP2005354	41.58	40.59	43.7	31.3	21.57	29.41	26.2	24.6	22.64	**	**	-
	NT2RP2005358	43.39	32.77	43.03	61.06	32.41	50.49	63.9	54.2	58.54	*		+
	NT2RP2005360	11.79	10	18.74	11.01	6.71	7.24	3.72	6.21	9.76			
	NT2RP2005378	29.24	21.46	29.8	11.99	9.12	7.48	6.73	9.68	15.33	**	*	-
	NT2RP2005391	8.87	18.64	25.48	15.21	9.76	9	4.05	7.34	6.04			
	NT2RP2005393	18.88	11.72	17.99	12.44	7.52	10.75	12.61	18.43	12.16			
55	NT2RP2005407	7.79	10.99	12.03	9.14	5.76	6.4	4.56	6.12	6.26	*		-
	NT2RP2005419	14.03	13.17	13.95	13.78	13.3	12.37	11.42	9.28	10.02	**		-
	NT2RP2005425	22.13	23.63	29.77	16.49	17.86	16.5	26.07	23.18	23.95	*		-

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	NT2RP2004410	39.53	65.74	46.52	39.11	16.74	16.36	21.28	23.23	25.45	*	-
	NT2RP2004412	14.74	10.36	20.72	26.06	1.96	8.17	8.89	4.91	6.36		
5	NT2RP2004414	8.26	7	7.21	7.75	3.36	2.32	8.71	6.3	5.47		
	NT2RP2004425	10.2	6.28	19.09	7.43	2.9	4.39	7.1	5.14	4.24		
	NT2RP2004447	7.9	4.2	15.6	7.23	1.76	4.51	4.43	8.04	6.17		
	NT2RP2004463	53.44	48.94	45.23	51.07	39.96	24.66	38.74	19.02	17.11	*	-
	NT2RP2004476	18.45	12.59	23.94	14.93	16.73	11.7	16.21	16.49	14.59		
	NT2RP2004488	10.21	10.37	13.58	8.53	5.46	7.65	8.97	6.18	4.15	*	*
10	NT2RP2004490	6.88	9.48	11.96	10.41	4.06	3.06	6.4	5.15	3.37		
	NT2RP2004495	59.84	40.86	52.11	47.08	29.44	39.52	53.9	45.74	40.44		
	NT2RP2004512	17.42	8.07	18.38	11.03	4.08	6.19	1.75	5.77	3.87	*	-
	NT2RP2004523	26.61	22.99	26.09	20.55	9.56	18.73	12.96	22.93	30.09		
	NT2RP2004524	21.22	18.85	29.36	17.14	12.55	10.77	11.27	10.59	14.51	*	-
	NT2RP2004536	39.58	17.04	37.57	26.29	26.05	14.38	14.91	28.86	19.32		
15	NT2RP2004538	145.51	113.79	134.59	95.27	89.81	81.41	118.33	103.7	132.33	*	-
	NT2RP2004548	21.33	25.87	20.88	16.6	11.36	9.17	14.62	15.2	15	*	**
	NT2RP2004551	8.55	11.05	18.95	8.38	5.63	4.59	5.77	5.04	6.54		
	NT2RP2004556	107.44	116.09	115.34	100.64	83.01	68.21	138.61	66.34	73.2	*	-
	NT2RP2004568	25.54	15.99	32.04	18.77	7.42	18.63	9.21	19.06	44.48		
	NT2RP2004580	28.48	22.84	30.46	23.04	12.01	15.3	24.34	30.63	24.49		
20	NT2RP2004585	104.19	75.97	102.59	102.16	96.71	82.97	64.51	65.78	63.06	*	-
	NT2RP2004587	7.64	7.71	10.73	6.38	2.94	4.03	2.08	3.66	11.64	*	-
	NT2RP2004594	9.49	13.7	13.53	10.94	10.06	5.96	7.18	7.17	14.25		
	NT2RP2004600	11.79	11.86	12.42	9.8	6.39	8.89	9.84	7.14	9.81	*	*
	NT2RP2004602	16.85	24.33	21.89	13.29	12.49	8.02	12.27	7.59	7.04	*	*
25	NT2RP2004606	282.85	252.35	268.41	181.43	249.55	245.26	257.06	189.04	158.13		
	NT2RP2004614	9.31	7.04	11.06	6.07	2.67	4.48	3.52	3.61	6.4	*	*
	NT2RP2004648	12.5	37.87	41.62	14.38	8.31	11.65	10.13	8.24	22.59		
	NT2RP2004655	27.02	40.5	41.28	26.01	23.69	13	4.9	10	13.87	**	-
	NT2RP2004664	12.3	12.14	15.06	9.64	6.17	5.52	2.65	4.39	2.54	*	**
	NT2RP2004670	9.81	11.79	9.6	6.17	5.86	2.45	3.26	4.14	2.42	*	**
30	NT2RP2004675	22.26	31.4	25.45	19.55	10.99	11.73	17.4	17	31.58	*	-
	NT2RP2004681	11.82	17.49	14.89	8.26	5.82	3.58	5.62	7.61	5.68	*	**
	NT2RP2004689	5.34	26.14	16.38	15.38	9.9	10.44	10.84	15.98	14.64		
	NT2RP2004709	38.28	22.19	35.75	22.32	11.28	19.42	7.33	20.31	16.27		
	NT2RP2004710	23.78	32.41	33.69	19.19	16.47	14	10.16	17.13	21.44	*	*
	NT2RP2004721	16.3	9.83	18.51	7.64	2.48	4.31	3.84	6.3	2.86	*	*
35	NT2RP2004736	12.51	12.2	14.69	12.94	8.12	6.27	5.93	10.36	6.34	*	-
	NT2RP2004743	18.86	18.19	16.11	11.52	9.54	6.56	6.31	8.72	9.61	**	**
	NT2RP2004750	59.2	56.12	53.52	47.6	43.74	38.38	59.22	63.07	62.81	**	-
	NT2RP2004755	27.3	33.26	34.09	17.59	21.76	19.12	17.05	12.23	8.96	**	**
	NT2RP2004767	16.81	19.28	18.07	14.61	10.15	10.32	11.83	8.67	9.96	*	**
	NT2RP2004768	39.27	17.22	28.44	49.02	23.23	30.46	23.63	20.93	17.76		
40	NT2RP2004775	10.4	6.35	15.81	7.59	3.37	4.94	1.86	3.35	3.03	*	-
	NT2RP2004791	35.3	37.93	43.25	33.95	20.1	16.04	17.4	29.11	27.1	*	-
	NT2RP2004794	54.46	52.25	59.56	37.23	24.39	16.04	14.21	17.04	17.83	*	**
	NT2RP2004795	16.82	13.04	16.28	12.97	8.52	3.96	6.33	6.26	8.64	**	-
	NT2RP2004799	16.76	14.17	14.52	9.03	7.47	5.55	2.73	2.76	3.58	**	**
	NT2RP2004802	42.99	46.43	41.16	46.78	39.32	30.76	28.53	29.92	38.26	*	-
45	NT2RP2004810	18.3	18.68	17.35	9.03	6.57	4.19	8.85	4.21	4.39	**	**
	NT2RP2004816	30.61	34.82	43.2	18.85	15.68	3.1	5.7	11.84	0.57	*	**
	NT2RP2004837	55.01	32.88	59.21	31.78	19.92	27.08	12.99	11.53	9.41	*	-
	NT2RP2004841	12.28	19.6	17.04	13.83	8.03	5.66	4.26	5.92	8.44	*	-
	NT2RP2004847	56	45.24	59.59	31.54	30.59	31.44	43.02	39.63	29.66	**	-
	NT2RP2004861	6.61	6.78	10.74	6.21	2.52	2.33	3.42	5.23	3.33		
50	NT2RP2004897	8.06	8.52	10.02	7.86	4.8	3.34	2.33	3.07	4.66	**	-
	NT2RP2004932	11.96	10.46	13.81	9.34	4.73	6.48	4.87	3.86	6	*	**
	NT2RP2004933	6.78	7.42	14.62	6.21	3.1	2.23	3.41	1.97	1.54	*	-
	NT2RP2004936	9.66	18.67	21.72	10.34	4.39	7.29	2.62	3.9	3.25	*	-
	NT2RP2004951	8.57	7.11	15.72	9.26	2.32	5.42	3.11	6.13	4.04		
	NT2RP2004959	12.5	13.69	22.18	6.53	5.09	6.63	3.92	5.67	5.43	*	*
55	NT2RP2004961	15.97	14.64	23.99	12.05	7.35	8.71	10.28	8.72	8.04	*	-
	NT2RP2004962	15.99	15.58	20.13	13.9	10.21	9.93	9.78	10.65	9.69	*	-
	NT2RP2004966	6.94	8.72	7.55	5.31	3.88	2.7	4.34	2.81	2.89	*	**

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	NT2RP2003952	8.61	6.4	10.47	5.99	3.42	3.14	3.11	4	4.05	*	*	-	-
	NT2RP2003968	14.45	22.39	24.81	11.6	25.12	6.38	6.57	9.01	14.39				
5	NT2RP2003976	33.5	36.02	32.11	28.15	26.29	22.25	25.49	23.11	18.45	*	**	-	-
	NT2RP2003981	10.38	7.73	15.47	7.98	3.66	7.9	3.93	9.3	6.45				
	NT2RP2003984	21.86	17.33	21.82	17.29	10	7.7	4.79	10.99	18.59				
	NT2RP2003986	29.04	20.99	42.5	17.22	10.08	14.95	20.31	27.16	19.47				
	NT2RP2003988	17.98	13.51	22.96	15.07	9.71	10.14	10.31	12.25	15.8				
	NT2RP2004013	76.9	66.49	68.74	48.94	44.57	34.09	28.41	25.31	32.75	**	**	-	-
10	NT2RP2004014	12.05	10.78	9.86	7.34	4.69	5.32	4.75	6.66	7.06	**	**	-	-
	NT2RP2004036	38.11	34.32	48.47	33.97	30.52	34.54	23.89	26.94	32.83				
	NT2RP2004041	15.68	9.6	13.65	12.24	6.88	6.29	9.46	7.39	4.47				
	NT2RP2004042	10.47	6.9	14.24	8.26	4.18	4.2	3.12	8.29	6.46				
	NT2RP2004049	50.08	29.35	58.86	54.03	47.21	47.08	23.48	30.4	35.04				
	NT2RP2004060	14.43	11.83	19.24	13.07	7.66	8.82	6.25	9.01	6.7	*		-	-
15	NT2RP2004066	9.22	7.65	15.12	7.14	3.11	3.37	2.62	3.51	3.15	*		-	-
	NT2RP2004069	13.07	11	19.34	9.29	5.59	3.82	4.85	6.14	4.78	*		-	-
	NT2RP2004076	7.23	8.97	8.94	6.15	5.2	2.62	3.55	3.66	4.01	*	**	-	-
	NT2RP2004080	16.76	19.32	21.22	10.81	7.94	4.93	7.71	6.49	6.54	**	**	-	-
	NT2RP2004081	12.13	9.06	12.95	7.67	4.89	5.2	4.76	5.29	3.41	*	**	-	-
	NT2RP2004098	14.04	9.1	18.6	7.17	4.55	6.94	5.74	6.96	7.08				
20	NT2RP2004108	15	13.48	22.62	10.06	6.21	6.99	5.4	7.73	10.17	*	*	-	-
	NT2RP2004124	8.21	9.75	17.19	8.66	5.64	4.42	4.13	6.1	2.63				
	NT2RP2004130	14.54	18.18	16.71	14.91	10.54	10.38	6.64	6.32	8.03	**		-	-
	NT2RP2004133	9.17	7.74	14.55	6.02	4.47	1.66	3.8	2.83	1.86	*		-	-
	NT2RP2004141	13.91	6.7	8.65	9.25	5.97	4.12	2.68	2.87	3.2	*		-	-
	NT2RP2004142	20.24	20.66	23.15	12.58	8.99	0.11	7.76	8.1	8.83	*	**	-	-
25	NT2RP2004152	12.94	17.34	28.22	13.04	12.79	7.85	8.66	8.13	4.02				
	NT2RP2004165	23.06	20.1	27.11	14.41	8.19	16.16	10.91	12.82	15.14	*	*	-	-
	NT2RP2004170	14.84	22.1	21.27	11.56	5.42	12.52	7.64	14.97	10.1	*		-	-
	NT2RP2004172	11.82	8.43	19.13	9.78	6.86	6.4	5.08	5.49	3.29				
	NT2RP2004176	16.04	24.68	27.17	14.08	13.85	12.56	15.42	21.18	13.53				
30	NT2RP2004179	11.06	11.88	18.99	6.12	3.68	3.62	3.12	4.46	3.04	*	*	-	-
	NT2RP2004187	10.62	14.3	17.05	10.83	6.49	6	6.3	3.3	5.65	*		-	-
	NT2RP2004190	11.44	13.08	11.65	11.65	16.98	5.99	14.27	4.91	8.09				
	NT2RP2004194	34.23	32.85	35.34	19.96	14.64	15.16	22.5	12.25	14.92	**	**	-	-
	NT2RP2004196	30.83	39.33	27.33	12.84	5.98	11.23	10.98	13.1	14.08	**	**	-	-
	NT2RP2004205	29.01	29.84	37.62	20.1	10.4	15.74	17.25	16.98	24.08	*	*	-	-
35	NT2RP2004207	8.99	13.88	16.49	8.05	7.02	4.72	8.6	7.78	25.16				
	NT2RP2004226	17.66	11.75	16.67	11.25	9.6	5.13	9.93	6.52	8.04	*		-	-
	NT2RP2004232	9.38	7.39	12.33	6.45	7.37	3.57	6.49	3.42	8.98				
	NT2RP2004239	9.12	12.33	10.22	7.52	5.19	3.55	2.71	5.07	3.29	*	**	-	-
	NT2RP2004240	34.67	45.84	33.56	20.39	18.64	6.67	14.35	6.97	4.81	*	**	-	-
	NT2RP2004242	15.64	15.83	16.65	14.76	9.29	9.76	10.22	6.01	6.98	**		-	-
40	NT2RP2004245	7.49	8.21	10.79	7.33	2.04	4.45	4.33	4.59	4.95	*		-	-
	NT2RP2004270	62.73	47.21	68.41	46.4	33.04	45.75	50.52	75.49	70.49				
	NT2RP2004300	12.51	8.53	13.1	8.54	9.07	5.04	6.79	9.38	7.79				
	NT2RP2004304	16.04	25.81	28.38	10.74	8.89	6.86	7.57	8.7	9.4	*	*	-	-
	NT2RP2004312	5.32	5.55	8.05	4.79	1.11	1.27	2.98	2.46	2.35	*		-	-
	NT2RP2004316	7.59	19.66	19.22	7.37	4.16	2.63	3.4	2.3	5.68	*		-	-
45	NT2RP2004321	11.35	10.01	12.72	6.18	1.74	2.16	3.62	2.37	4.34	**	**	-	-
	NT2RP2004336	6.56	6.01	5.01	7.15	3.83	3.55	3.86	1.85	3.64	*		-	-
	NT2RP2004339	86.87	54.65	64.97	50.21	27.02	70.75	73.53	111.22	98.32				
	NT2RP2004347	9.56	12.29	15.81	9.78	3.94	5.75	4.47	6.18	9.27				
	NT2RP2004364	17.82	15.18	19.5	11.98	7.62	12.3	9.63	15.21	13.78	*		-	-
	NT2RP2004365	16.43	16.15	23.6	8.67	8.61	4.88	6.23	5.81	4.59	*	**	-	-
50	NT2RP2004366	6.71	10.58	9.59	7.66	3.53	5.17	5.46	3.22	3.29	*		-	-
	NT2RP2004373	12.26	11.21	11.74	7.26	2.71	4.58	3.35	3.56	4.51	**	**	-	-
	NT2RP2004375	63.47	77.52	81.72	36.96	16.48	21.89	34.46	29.81	18.56	**	**	-	-
	NT2RP2004389	11.31	16.38	12.26	9.91	2.21	4.95	5.82	4.38	4.02	*	**	-	-
	NT2RP2004392	57.13	31.73	43.78	20.38	16.25	19.24	10.86	16.27	14.58	*	*	-	-
	NT2RP2004396	13.42	9.2	11.06	9.69	3.89	7.39	5.67	7.43	7.76	*		-	-
55	NT2RP2004399	25.77	19.74	26.11	15.91	12.56	15.11	9.21	12.21	10.01	*	**	-	-
	NT2RP2004400	10.29	9.25	9.33	9.9	4.17	8.44	7.31	8.61	9.43	*		-	-
	NT2RP2004404	142.91	126.47	123.47	87.01	85.43	109.26	171.94	127.81	174.94	*		-	-

Table 430

	NT2RP2003506	18.91	15.11	21.41	10.64	13.4	7.04	4.15	4.83	3.94	*	**	-	-
	NT2RP2003511	12.05	3.9	4.64	8.42	8.48	3	2.22	2.49	1.23			-	-
5	NT2RP2003513	5.02	4.48	4.09	4.53	1.64	2.04	2.58	1.55	3.7	*		-	-
	NT2RP2003517	3.09	1.94	2.24	4.01	1.27	1.81	1.81	0.62	1.53			-	-
	NT2RP2003522	5.79	5.52	8.11	4.61	3.87	1.76	1.31	4.96	5.44			-	-
	NT2RP2003525	12.34	13.08	16.44	6.02	5.34	4.08	6.04	6.6	5.02	**	**	-	-
	NT2RP2003533	88.01	167.21	169.95	95.54	157.31	89.62	178.82	120.18	151.89			-	-
	NT2RP2003541	3.67	3.34	5.25	3.35	1.69	1.26	1.11	0.78	0.47		**	-	-
10	NT2RP2003543	6.02	7.31	6.9	4.81	3.89	1.42	3.96	1.71	3.06	*	**	-	-
	NT2RP2003545	16.66	28.91	37	10.57	5.83	7.38	9	7.02	10.57	*	*	-	-
	NT2RP2003559	24.73	30.59	37.42	18.05	13.28	13.04	17.81	5.8	6.39	*	*	-	-
	NT2RP2003564	4.21	3.5	2.84	5.11	2.21	2.76	1.47	1.11	1.54	**		-	-
	NT2RP2003565	5.56	6.05	9.75	8.3	2.97	4.12	3.13	4.62	3.34			-	-
	NT2RP2003567	52.47	49.61	31.71	25.62	16.56	16.12	8.17	19.23	14.83	*	*	-	-
15	NT2RP2003575	228.18	422	264.34	79.35	234.74	114.07	248.45	399.29	319.2			-	-
	NT2RP2003576	12.51	13.48	14.93	11.41	10.35	7.47	9.35	12.38	11.58	*		-	-
	NT2RP2003579	6.35	14.72	8.69	5.77	5	3.85	5.14	3.96	3.9			-	-
	NT2RP2003581	5.18	8.39	5.74	5.34	1.72	1.57	1.66	0.89	1.73	**		-	-
	NT2RP2003587	5.23	4.31	3.93	4.85	1.48	2.81	2.91	3.12	2.6	*		-	-
20	NT2RP2003590	10.26	7.56	7.31	8.22	4.4	10.25	8.81	5.02	6.24			-	-
	NT2RP2003593	30.36	26.83	29	16.89	17.04	17.86	10.43	21.71	15.35	**	*	-	-
	NT2RP2003596	9.32	8.79	11.25	13.55	9.73	9.49	8.79	14.74	9.47			-	-
	NT2RP2003599	23.27	13.17	35	8.05	27.81	6.94	3.86	4	10.53			-	-
	NT2RP2003600	19.86	30.4	31.86	22.5	9.14	19.7	15.99	13.53	9.08	*	*	-	-
	NT2RP2003604	5.88	9.89	7.47	5.4	1.89	1.85	3.39	3.16	2.76	*	*	-	-
25	NT2RP2003629	8.19	7.7	5.26	5.58	3.64	2.51	3.1	3.51	2.87	*	*	-	-
	NT2RP2003630	4.6	3.1	1.8	4.67	0.47	1.48	1.43	1.9	2.02			-	-
	NT2RP2003643	14.26	14.44	12.75	14.01	7.67	9.12	19.42	10.04	13.07			-	-
	NT2RP2003655	7.62	6.24	7.73	7.33	2.44	6.8	2.97	5.19	4.57	*		-	-
	NT2RP2003664	8.96	13.57	11.03	11.23	17.02	7.98	3.52	3.83	3.73	**		-	-
	NT2RP2003668	5.83	2.7	3.35	6.14	1.92	4.61	2.43	9.15	4.25			-	-
30	NT2RP2003687	2.45	2.75	3.32	4.63	0.64	2.77	0.71	2.3	0.91			-	-
	NT2RP2003691	2.86	0.41	1.55	4.29	1.06	2.07	1.69	2.35	1.51			-	-
	NT2RP2003702	7.99	4.44	8.03	9.77	1.89	6.56	5.26	5.52	3.42			-	-
	NT2RP2003704	6.11	9.87	8.33	7.33	6.55	3.79	6.43	2.95	5.24			-	-
	NT2RP2003706	7.36	9.13	9.09	6.95	3.88	3.8	6.16	3.16	4.56	*	*	-	-
	NT2RP2003713	10.55	5.71	14.4	8.75	4.92	4.56	5.34	3.61	4.08			-	-
35	NT2RP2003714	8.27	4.37	12.63	8.63	2.38	6.34	8.02	5.56	6.23			-	-
	NT2RP2003727	13.72	9.91	16.65	14.5	8.49	5.28	4.4	3.92	5.54	*		-	-
	NT2RP2003737	5.45	5.17	12.85	7.76	3.42	3.25	3.61	5.76	7.42			-	-
	NT2RP2003751	4.58	1.12	4.62	4.58	0.42	0.71	2.16	1.87	0.69			-	-
	NT2RP2003760	5.71	6.37	7.89	7.16	2.44	3.65	5.58	4.39	4.01			-	-
	NT2RP2003764	4.38	3.82	3.93	5.42	1.13	2.52	3.39	1.74	2.45	*		-	-
40	NT2RP2003769	20.81	10.06	18.85	16.88	9.37	7.83	5.73	10.32	7.17			-	-
	NT2RP2003770	46.73	23.49	36.27	27.58	17.68	15.16	13.53	17.82	25.92			-	-
	NT2RP2003777	15.17	10.4	15.63	9.4	5.35	5.3	5.51	8.22	8.17	*	*	-	-
	NT2RP2003781	85.09	53.2	80.23	44.9	32.89	44.6	88.34	75.84	75.38	*		-	-
	NT2RP2003785	11.62	18.04	20.61	13.78	15.2	6.96	6.32	9.84	5.9	*	*	-	-
	NT2RP2003793	16.4	16.91	28.84	17.84	16.4	19.5	8.6	5.86	10.58	*		-	-
45	NT2RP2003806	22.61	30.87	30.72	19.98	15.24	17.95	21.46	20.85	31.8	*		-	-
	NT2RP2003825	121.72	69.01	127.36	108.65	103.08	122.02	106.86	101.92	114.26			-	-
	NT2RP2003840	11.66	9.2	14.49	9.94	6.01	4.99	5.06	8.29	6.32	*		-	-
	NT2RP2003857	12.37	17.33	20.12	13.2	12.38	5.39	3.92	13.14	19.37			-	-
	NT2RP2003859	19.27	8.66	21.11	12.54	7.05	8.25	8.14	9.73	13.41			-	-
	NT2RP2003871	24.21	21.74	19.35	11.95	9.6	10.67	5.81	8.41	8.75	**	**	-	-
50	NT2RP2003876	10.53	11.98	18.66	19.02	25.1	11.05	7.68	10.36	8.8			-	-
	NT2RP2003878	8.32	9.1	10.32	8.36	3.76	3.6	4.02	3.95	3.32	**		-	-
	NT2RP2003885	6.08	8.46	6.77	5.72	4.33	2.2	4.11	3.32	4.21	*		-	-
	NT2RP2003898	16.47	22.78	20.73	12.55	8.77	11.94	7.99	8.27	7.33	*	**	-	-
	NT2RP2003902	21.99	10.54	20.38	16.69	8	10.58	8.28	15.68	15.89			-	-
	NT2RP2003912	35.16	73.65	32.26	21.86	23.82	12.49	13.87	18.46	19.33			-	-
55	NT2RP2003931	16.31	15.84	40.63	15.98	12.61	14.09	15.19	20.43	45.95			-	-
	NT2RP2003940	26.29	28.19	28.81	19.73	11.64	14	20.12	28.2	22.66	**		-	-
	NT2RP2003950	10.42	7.47	13.08	10.4	5.75	5.62	6.83	8.37	6.11			-	-

Table 429

	NT2RP2003125	148.62	154.21	165.76	136.93	87.76	119.3	202.39	170.18	181.03				
	NT2RP2003127	8.19	13.35	12.24	9.29	4.08	4.24	7.52	6.27	6.75				
5	NT2RP2003129	5.37	4.98	2.74	4.8	1.83	1.5	2.73	2.54	2.23				
	NT2RP2003137	9.57	8.81	8.47	10.46	3.57	7.13	10.02	8.73	11.39				
	NT2RP2003138	289.53	193.02	256.72	147.84	144.9	131.2	108.58	156.98	150.94	*	*	-	-
	NT2RP2003146	1.24	1.09	7.94	3.13	0.31	1.29	1.86	1.99	1.77				
	NT2RP2003148	8.05	4.25	16.47	5.03	5.56	4.21	3.03	3.24	3.73				
10	NT2RP2003150	10.24	5.66	13.18	8.31	3.11	3.88	5.08	3.43	3.42				
	NT2RP2003157	3.48	4.11	8.05	5.53	2.9	2.53	3.08	2.08	2.29				
	NT2RP2003158	154.65	121	143.84	65.15	86.73	95.34	71.82	47.48	71.44	*	**	-	-
	NT2RP2003161	7.49	10.19	13.16	11.02	5.25	6.57	6.09	5.72	4.72	*		-	
	NT2RP2003164	3.55	3.69	5.64	7.04	3.38	4.04	7.46	5.26	5.34				
	NT2RP2003165	12.93	8.52	16.52	10.68	5.85	8.09	3.19	7.18	6.13				
15	NT2RP2003177	20.41	17.38	32.83	19.51	10.51	7.06	22.73	21.74	21.1				
	NT2RP2003179	13.75	8.83	17.7	6.83	5.73	5.2	6.82	13.09	8.84	*		-	
	NT2RP2003194	60.36	55.91	80.01	51.34	60.6	65.85	42.59	38.48	56.22				
	NT2RP2003206	4.24	5.6	7.88	6.81	4.57	3.96	3.75	3.9	1.77				
	NT2RP2003210	9.9	9.36	13.05	7.55	5.04	4.08	4.47	4.65	4.08	*	**	-	-
	NT2RP2003227	13.08	17.16	19.16	9.3	7.87	5.3	8.38	6.37	6.65	*	**	-	-
20	NT2RP2003228	16.97	20.02	17.73	19.68	23.22	23.38	16.96	9.41	24.11				
	NT2RP2003230	8.37	4.38	11.24	9.26	4.24	4.38	5.34	6.55	20.33				
	NT2RP2003231	19.49	17.75	28.22	19.04	13.83	13.5	4.32	15.06	15.52				
	NT2RP2003237	7.53	6.29	14.94	7.53	4.21	4.44	2.72	5.12	9.5				
	NT2RP2003239	14.69	12.62	13.82	13.64	7.63	7.59	6.01	11.3	7.74	*		-	
	NT2RP2003243	18.5	16.09	21.39	14.68	13.35	11.44	13.72	14.6	10.6	*	*	-	-
25	NT2RP2003265	4.16	3.24	4.88	3.11	1.7	1.76	2.29	2.88	1.56	*	*	-	-
	NT2RP2003267	60.51	57	71.61	43.05	41.52	45.26	62.57	46.27	55.1	*		-	
	NT2RP2003272	18.93	12.31	21.16	15.62	7.25	9.68	9.71	5.82	7.07	*	*	-	-
	NT2RP2003277	51.75	42.05	60.1	30.74	24.84	22.69	27.09	34.52	27.77	*	*	-	-
	NT2RP2003280	28.6	31.79	33.98	27.16	26.4	22.37	7.26	22.23	17.9	*	*	-	-
	NT2RP2003286	5.27	5.54	11.77	5.48	3.75	3.02	2.08	4.89	2.85				
30	NT2RP2003293	449.62	544.88	531.61	480.84	613.42	591.27	492.68	295.17	642.99				
	NT2RP2003295	31.05	28.41	29.77	21.7	14.01	19.1	19.18	18.46	19.51	**	**	-	-
	NT2RP2003297	16.92	17.67	19.83	9.38	11.73	7.84	3.24	6.26	6.08	**	**	-	-
	NT2RP2003300	2.42	4.36	4.19	5.32	2.8	1.56	2.33	1.69	1.31				
	NT2RP2003302	4.27	4.34	3.48	7.02	4.29	2.92	3.56	4.67	4.2				
	NT2RP2003307	17.52	15.84	29.67	14.93	8.5	10.4	16.58	17.1	22.41				
35	NT2RP2003308	7.26	8.66	14.15	6.9	4.51	4.44	3.22	4.76	6.65				
	NT2RP2003311	29.64	20.85	41.04	22.79	16.59	15.7	26.2	28.58	26.19				
	NT2RP2003329	11.36	11.27	19.3	9.84	10.72	8.3	6.72	6.84	5.55	*		-	
	NT2RP2003339	44.06	51.39	68.66	63.28	61.45	41.92	35.03	31.79	30.76	*		-	
	NT2RP2003345	25.41	28.87	30.12	15.65	24.89	17.7	13.62	14.02	13.58	*	**	-	-
	NT2RP2003347	8.25	20.81	11.17	7.75	7.09	4.08	5.16	5.37	4.53				
40	NT2RP2003357	15.71	9.07	15.92	25.01	12.88	13.97	19.6	13.55	17.34				
	NT2RP2003359	28.62	21.24	26	8.89	11.95	10.09	4.71	10.13	9.11	**	**	-	-
	NT2RP2003383	14.02	11.63	23.31	10.33	7.37	5.29	4.94	12.01	8.63				
	NT2RP2003390	5.02	4.43	9.92	6.51	4.16	1.23	2.19	2.28	1.48				
	NT2RP2003391	7.9	10.18	14.85	7.86	4.63	3.15	3.6	3.53	4.66	*		-	
	NT2RP2003393	17.88	10.78	13.47	5.99	12.67	3.97	3.38	3.45	4.48	**		-	
45	NT2RP2003394	86	76.58	96.35	85.96	75.14	69.1	44.7	55.38	51.85	**		-	
	NT2RP2003401	6.44	4.6	7.19	7.26	3.17	2.2	2.64	2.79	2.35	*		-	
	NT2RP2003403	18.73	19.58	18.59	16.65	31.07	15.94	9.34	7.98	7.02	**		-	
	NT2RP2003433	8.13	7.64	13.92	4.56	2.36	2.9	0.77	3.05	2.35	*	*	-	-
	NT2RP2003445	11.02	10.94	17.55	8.15	5.02	3.19	3.72	5.83	4.71	*	*	-	-
	NT2RP2003446	7.38	5.87	15.32	8.9	3.31	1.64	2.84	3.34	2.61				
50	NT2RP2003456	14.58	16.09	17.24	7.76	9.78	4.68	6.41	6.61	4.94	**	**	-	-
	NT2RP2003466	163.89	136.64	191.26	110.44	140.48	151.43	83.63	87.04	121.39	*		-	
	NT2RP2003469	5.44	3.7	5.03	4.74	2.31	1.35	2.28	1.42	1.62	**		-	
	NT2RP2003470	4.83	5.55	5.49	5.65	3.19	3.4	4.42	1.43	3				
	NT2RP2003471	12.29	11.22	13.81	11.17	23.08	8.88	5.31	3.9	2.98	**		-	
	NT2RP2003480	133.45	297.88	151.87	131.22	166.23	117.7	104.84	202.75	202.47				
55	NT2RP2003495	2.81	3.4	10.31	5.19	1.55	2.41	0.68	1.49	0.7				
	NT2RP2003499	5.25	3.3	9.47	3.94	2.15	2.35	2.44	6.25	5.89				
	NT2RP2003505	1.66	1.94	6.72	4.41	2.46	0.11	0.21	3	0				

Table 428

	NT2RP2002673	103.76	111.24	114.16	70.07	45.09	85.59	92.44	85.88	91.14	*	**	-	-
	NT2RP2002674	28.24	27.69	34.14	21.06	20.78	16.49	24.37	22.8	22.81	*	*	-	-
5	NT2RP2002686	20.76	27.04	31.72	23.26	22.04	15.33	8.73	8.68	8.13	**		-	-
	NT2RP2002688	13.92	12.24	10	14.56	10.69	8.28	14.43	8.67	10.91			-	-
	NT2RP2002695	19.25	14.55	23.17	14.75	11.8	15.12	11.5	15.77	21.62			-	-
	NT2RP2002701	4.75	5.4	12.91	6.52	3.02	5.45	3.43	3.42	5.6			-	-
	NT2RP2002706	14.53	15.48	22.22	9.32	11.09	8.76	5.34	11.12	9.16	*	*	-	-
10	NT2RP2002710	30.61	29.84	37.75	21.5	19.35	13.67	11.45	15.23	13.68	*	**	-	-
	NT2RP2002721	7.29	6.84	7.35	5.81	4.84	2.96	3.61	3.96	1.62	*	**	-	-
	NT2RP2002727	6.54	9.38	10.59	7.9	7.44	4.24	4.08	5.05	3.6	*		-	-
	NT2RP2002734	48.13	41.34	40.09	29.15	43.09	31.56	39.87	34.2	33.11			-	-
	NT2RP2002736	9.71	9.47	8.91	9.58	7.4	9.48	7.64	7.21	5.83	*		-	-
	NT2RP2002740	19.4	13.22	23.87	11.87	7.31	11.43	6.19	11.24	13.39			-	-
15	NT2RP2002741	23.72	17.52	33.51	16.7	12.26	14.24	12.67	22.55	22.7			-	-
	NT2RP2002750	63.6	50.03	64.42	39.1	24.44	35.13	36.13	36.29	30.98	*	**	-	-
	NT2RP2002752	10.52	9.47	15.12	8.58	15.15	3.88	2.8	5.03	3.54	*		-	-
	NT2RP2002753	12.15	11.11	17.25	9.74	6.82	4.55	5.04	8.44	4.88	*		-	-
	NT2RP2002760	73.48	81.4	101.38	101.02	88.73	62.2	63.54	64.02	58.41			-	-
	NT2RP2002769	92.25	85.07	85.98	43.44	97.23	54.77	13.8	14.78	19.89	**		-	-
20	NT2RP2002778	10.58	7.74	8.45	8.8	4.88	4.14	4.05	4.2	5.19	**		-	-
	NT2RP2002791	21.96	17.69	24.67	15.97	10.79	19.44	8.98	15.96	17.2			-	-
	NT2RP2002800	57.56	44.66	67.26	27.41	29.24	35	38.77	66.85	61.35	*		-	-
	NT2RP2002805	10.55	19.25	27.64	13.98	23.59	9.88	7.4	10.07	7.75			-	-
	NT2RP2002811	13.01	13.43	18.61	12.19	7.67	8.64	7.15	10.12	7.7	*		-	-
	NT2RP2002824	16.63	12.33	20.93	15.22	10.57	11.14	9.08	9.37	7.94	*		-	-
25	NT2RP2002839	7.61	6.4	5.94	5.9	4.06	3.3	9.77	3.03	3.13			-	-
	NT2RP2002845	12.27	9.25	11.68	10.47	8.06	5	6.84	6.96	4.06	*		-	-
	NT2RP2002857	13.78	16.15	14.57	14.65	30.41	11.09	9.34	6.45	5.36	**		-	-
	NT2RP2002862	39.8	48.81	41.43	30.51	20.27	24.76	25.07	37.41	27.22	*	*	-	-
	NT2RP2002880	68.2	65.99	77.82	51.13	34.35	43.94	48.02	65.53	66.61	*		-	-
	NT2RP2002885	6.6	7.46	15.43	8.27	7.08	4.47	4.01	3.85	3.27			-	-
30	NT2RP2002891	7.27	7.43	13.2	9.04	9.22	4.71	4.88	7.24	5.8			-	-
	NT2RP2002907	27.84	27.28	39.25	27.93	20.22	19.34	24.71	20.15	30.92			-	-
	NT2RP2002925	8.14	12.06	8.92	6.22	5.24	3.39	2.57	2.53	2.32	*	**	-	-
	NT2RP2002927	11.04	11.53	9.75	8.5	11.71	5	4.88	3.47	4.87	**		-	-
	NT2RP2002928	14.93	19.65	15.07	19.82	17.04	17.08	11.05	6.88	7.13	*		-	-
	NT2RP2002929	34.38	37.86	32.48	20.14	17.6	22.45	18.17	27.25	31.67	**		-	-
35	NT2RP2002934	25.44	20.87	30.69	18.19	13.85	16.46	17.37	22.76	21.39	*		-	-
	NT2RP2002939	16.49	24.31	21.25	14.42	15.1	11.79	7.22	8.55	7.99	*	**	-	-
	NT2RP2002942	60.99	80.16	86.43	63.29	72.69	51.11	69.34	65.39	67.68			-	-
	NT2RP2002954	14.37	21.3	26.97	15.75	10.45	11.22	8.24	9.15	9.35	*		-	-
	NT2RP2002959	11.9	10.85	14.73	7.3	8.01	4.57	5.17	3.69	3.58	*	**	-	-
40	NT2RP2002974	27.97	23.95	30.02	18.63	11.35	13.46	21.06	14.1	16.2	*	*	-	-
	NT2RP2002976	33.37	27.08	30.74	41.03	19.52	40.87	48.18	35.03	46.65	*		-	-
	NT2RP2002979	34.44	28.83	35.03	24.79	13.64	25.53	28.21	33.42	38.67			-	-
	NT2RP2002980	56.79	52.53	62.74	67.66	61.62	49.88	36.91	45.4	40.52	*		-	-
	NT2RP2002986	8.8	8.37	8.86	6.06	9.31	3.85	4.83	5.06	6.03	**		-	-
	NT2RP2002987	103.6	100.12	86.59	73.66	56.64	94.83	99.25	107.73	109.15			-	-
45	NT2RP2002988	13.55	24.45	23.43	8.39	5.16	5.67	6.67	7.13	6.64	*	*	-	-
	NT2RP2002993	30.07	31.12	32.86	20.82	20.33	19.14	25.73	28.68	29.96	**		-	-
	NT2RP2003000	23.19	27.52	36.47	14.21	13.97	11.57	16.09	13.68	17.86	*	*	-	-
	NT2RP2003008	20.38	21.66	17.68	30.65	18.58	30.92	20.83	11.64	23.71			-	-
	NT2RP2003020	8.4	9.59	8.76	9.59	3.53	4.92	4.04	6.43	6.19	*		-	-
	NT2RP2003032	16.62	21.12	22.97	11.47	10.11	7.97	7.7	9.72	7.64	**	**	-	-
50	NT2RP2003034	9.72	12.06	10.53	7.37	15.19	4.72	7.75	4.61	8.9			-	-
	NT2RP2003042	7.27	6.68	12.91	7.34	5.07	3.54	6.69	5.74	4.38			-	-
	NT2RP2003050	41.14	39.77	34.47	22.7	14.89	22.18	30.79	33.73	32.33	**	*	-	-
	NT2RP2003080	13.62	10.73	6.38	6.73	19.61	3.28	5.01	3.28	4.49			-	-
	NT2RP2003073	11.88	8.02	5.9	7.53	3.76	4.73	2.61	3.98	5.14			-	-
	NT2RP2003099	15.8	18.78	12.88	19.36	10.86	16.96	13.15	11.19	10.41			-	-
	NT2RP2003108	16.99	10.19	16.94	12.41	6.81	13.81	6.41	10.09	7.95			-	-
55	NT2RP2003115	140.52	119.44	125.07	149.82	106.25	154.27	145.7	152.43	149.71	*		-	-
	NT2RP2003117	8.58	4.12	8.6	7.48	2.66	5.97	6.92	5.37	6.56			-	-
	NT2RP2003121	15.37	11.96	21.6	10.75	8.2	9.97	7.29	7.74	8.68	*		-	-

Table 427

	NT2RP2002188	222.52	243.75	126.54	151.89	124.73	74.73	64.57	82.95	98.67	*	-
	NT2RP2002192	66.59	140.59	123.86	61.74	74.83	59.86	111.82	74.79	70.77		
5	NT2RP2002193	9.92	9.38	5.54	5.64	8.2	3.35	4.38	2.62	2.3	*	-
	NT2RP2002208	24.07	21.97	26.87	14.79	10.85	11.7	14.39	13.9	16.1	**	-
	NT2RP2002219	67.68	67.68	54.54	42.26	45.01	39.83	12.23	52.49	39.63	*	-
	NT2RP2002231	7.62	8.91	9.95	5.33	3.41	3.41	3.43	5.73	5.85	**	-
	NT2RP2002232	82.24	77.63	81.5	78.38	94.62	65.26	53.25	57.27	56.31	**	-
	NT2RP2002235	28.16	33.42	30.04	28.68	25.16	22.07	20.58	21.22	23.01	**	-
10	NT2RP2002239	46.02	63.63	67.29	21.58	16.63	25.15	34.54	22.68	19.45	**	-
	NT2RP2002252	7.05	9.07	9.23	8.29	3.79	3.9	4.57	3.45	2.91	**	-
	NT2RP2002256	27.97	26.83	27.09	15.84	11.06	11.47	26.86	22.34	28.47	**	-
	NT2RP2002257	7.18	9.4	7.52	6.89	3.78	4.59	5.56	4.76	6.16	*	-
	NT2RP2002259	18.53	13.43	11.26	24.93	14.6	14.24	10.77	18.34	15.94		
	NT2RP2002264	26.23	33.28	22.8	12.91	13.58	12.04	9.7	17.76	13.48	**	-
15	NT2RP2002267	32.17	30.65	32.28	23.04	23.01	26.57	18.83	21.9	22.12	**	-
	NT2RP2002270	24.56	42.02	39.93	12.58	23.51	8.23	20.01	17.31	11.5	*	-
	NT2RP2002281	77.89	96.13	97.81	150.54	78.9	96.99	77.36	60.86	55.61	*	-
	NT2RP2002288	21.39	25.5	23.86	18.7	13.7	14.18	19.97	22.13	17.3	*	-
	NT2RP2002292	54.6	51.47	64.64	33.71	24.08	25.61	33.57	27.11	23.57	**	-
	NT2RP2002299	38.27	50.42	61.77	19.48	10.3	12.83	10.02	9.38	6.36	**	-
20	NT2RP2002304	11.94	6.17	8.22	10.44	3.85	9.51	5.62	11.16	9.83		-
	NT2RP2002312	27.46	21.25	33.99	24.99	23.29	27.25	22.52	32.61	25.31		-
	NT2RP2002316	51.29	40.93	55.81	26.92	27.74	16.77	8.08	18.25	13.09	*	-
	NT2RP2002325	14.79	13.26	29.38	13.29	7.49	7.83	6.29	6.54	6.53	*	-
	NT2RP2002333	37.71	73.03	60.93	25.88	22.98	32.97	22.31	24.03	28.79	*	-
	NT2RP2002371	18.55	14.9	20.81	15.66	7.83	11.84	19.13	18.46	18.7		-
25	NT2RP2002373	7.7	6.37	4.6	6.91	3.79	4.83	4.84	5.13	5.14		-
	NT2RP2002381	9.55	13.3	13.26	6.59	3.77	3.7	6.54	4.52	6.36	**	-
	NT2RP2002385	0.45	2.78	2.19	4.9	1.81	3.5	0.24	0.66	1.05		-
	NT2RP2002394	32.72	28.71	26.62	28.1	25	26.16	15.67	15.8	20.33	**	-
	NT2RP2002408	6.97	5.52	5.11	6.31	2.85	4.31	2.1	5.36	5.24		-
	NT2RP2002409	7.43	7.51	7.75	10.28	4.99	7.17	4.61	5.21	4.74	**	-
30	NT2RP2002424	13.74	18.21	4.21	14.47	4.22	10.67	7.29	17.72	5.01		-
	NT2RP2002426	31.08	22.01	26.45	29.33	10.8	28.89	38.27	34.35	51.45		-
	NT2RP2002429	5.25	5.45	6.58	6.82	3.95	4.28	5.54	2.96	5.77		-
	NT2RP2002437	5.68	8.15	5.2	8.99	3.78	4.37	5.91	5.66	5.8		-
	NT2RP2002439	23.83	11.85	28.63	30	21.32	18.74	18.69	27.87	21.85		-
	NT2RP2002442	4.21	2.48	10.06	7.13	2.17	3.83	3.89	4.18	3.7		-
35	NT2RP2002457	5.91	3.86	9.43	6.08	1.82	4.35	4.32	3.38	4.15		-
	NT2RP2002464	14.04	9.04	19.39	8.26	2.91	3.02	6.2	7.3	8.31		-
	NT2RP2002475	18.47	16.05	30.87	26.73	13.63	23.54	27.01	26.72	27.62		-
	NT2RP2002479	4.38	4.25	7.68	9.08	3.28	4.87	3.36	7.1	3.51		-
	NT2RP2002487	8.46	11.73	16.26	11.78	4.98	7.15	5.71	5.33	8.37		-
40	NT2RP2002496	12.42	14.56	14.54	14.3	15.74	13.48	11.49	11.5	14.04		-
	NT2RP2002503	38.54	26.71	38.28	36.97	26.13	31.51	31.04	37.76	40.27		-
	NT2RP2002504	10.58	11.45	20.66	9.47	10.7	9.29	7	7.64	9.65		-
	NT2RP2002510	35.82	27.78	43.7	25.78	25.92	21.7	29.92	29.43	28.5		-
	NT2RP2002520	21.45	19.39	27.79	16.61	13.24	8.38	14.06	12.04	8.59	*	-
	NT2RP2002527	36.1	23.32	35.4	20.76	25.32	20.27	27.42	26.23	15.63		-
45	NT2RP2002533	13.17	14.75	23.11	11.93	9.58	7.87	8.62	7.96	9		-
	NT2RP2002537	13.08	12.39	14.79	10.65	13.69	8	8.5	6.59	7.09	**	-
	NT2RP2002542	68.44	80.08	72.33	74.6	46.95	45.75	49.02	39.47	37.92	**	-
	NT2RP2002546	14.88	18.96	17.6	15.64	9.68	8.83	6.06	11.83	13.87		-
	NT2RP2002549	14.96	11.64	16.89	10.43	8.76	6.03	5.02	9.52	9.5	*	-
	NT2RP2002584	39.56	26.21	34.95	20.11	17.81	17.77	18.21	24.98	19.23	*	-
50	NT2RP2002591	42.59	28.9	37.43	32.97	30.53	25.24	25	30.71	25.2		-
	NT2RP2002595	5.68	8	12.85	8.34	7.39	5.51	5.35	4.75	4.31		-
	NT2RP2002602	54.63	55.87	75.68	59.64	57.23	46.71	59.72	54.59	54.85		-
	NT2RP2002606	12.52	13.19	14.5	13.95	17.61	9.8	13.98	12.61	14.58		-
	NT2RP2002609	11.6	17.33	13.31	20.12	19.43	14.55	10.05	12.64	13.06		-
	NT2RP2002618	14	10.22	19.04	13.28	7.51	9.59	7.56	12.06	11.33		-
55	NT2RP2002621	11.85	9.77	15.54	8.04	5.31	5.01	3.35	5.97	7.64	*	-
	NT2RP2002643	11.04	8.19	16.62	9.29	6.05	6.59	3.63	6.53	4.9		-
	NT2RP2002672	6.88	6.19	9.33	7.99	5.59	4.36	3.64	6.54	3.29		-

Table 426

	NT2RP2001769	74.25	64.35	76.05	49.19	51.48	48.88	16.14	29.26	21.95	**	**	-	-
	NT2RP2001784	6.1	6.64	9.79	6.92	5.74	5.64	4.78	4.07	4.19				
5	NT2RP2001805	18.64	23.08	23.59	18.75	15.28	15.88	12.77	15.01	19.71				
	NT2RP2001813	12.2	15.41	15.49	8.88	4.21	6.14	9.29	6.92	9.66	*	*	-	-
	NT2RP2001817	50.4	58.09	53.81	41.79	32.65	39.63	54.02	42.75	56.58	*		-	
	NT2RP2001818	7.27	4.11	3.7	5.91	1.54	5.16	5.71	4.24	6.18				
	NT2RP2001837	24.91	17.75	31.06	23.35	23.21	19.49	21.43	23.69	22.4				
10	NT2RP2001839	16.58	14.52	27.11	17.12	15.87	19.03	13.8	13.73	15.93				
	NT2RP2001861	19.37	18.6	26.95	11.83	7.45	7.51	9.29	9.04	10.22	*	*	-	-
	NT2RP2001869	8.35	8.88	14.85	9.01	6.82	4.27	3.19	6.52	4.2				
	NT2RP2001876	52.07	48.1	60.43	41.56	38.63	33.22	53.55	54.23	52.55	*		-	
	NT2RP2001878	10.64	8.11	16.99	8.71	6.81	6.18	8.15	6.26	6.36				
	NT2RP2001881	11.51	12.01	17.12	11.39	8.77	9.36	10.8	14.93	9.65				
	NT2RP2001883	15.67	6.93	15.2	18.32	7.55	11.74	13.41	14.76	12.05				
15	NT2RP2001884	1443.8	767.12	607.01	701.8	366.77	733.69	199.74	492.38	1929.8				
	NT2RP2001885	86.63	81.18	119.19	86.34	68.67	72.27	80.49	93.38	87.15				
	NT2RP2001898	13.46	6.9	14.73	6.96	6.72	3.84	5.67	4.43	5				
	NT2RP2001900	20.73	14.58	24.69	18.79	8.43	10.74	17.28	16.1	17.71				
	NT2RP2001903	11.05	14.47	16.38	12.77	7.63	7.86	11	14.28	12.43				
20	NT2RP2001907	9.79	12.89	12.82	8.11	6.3	4.57	8.52	7.33	7.63	*	*	-	-
	NT2RP2001915	17.49	19.44	20.82	11.45	7.41	6.04	7.94	7.59	6.62	**	**	-	-
	NT2RP2001921	13.67	9.05	20.66	11.35	6.61	9.21	6.21	10.48	17.99				
	NT2RP2001926	34.47	22.9	35.27	22.68	13.7	11.83	10.15	26.13	26.8	*		-	
	NT2RP2001933	36.39	16.3	35.12	23.35	20.81	17.87	10.85	17.5	22.37				
	NT2RP2001936	11.42	9.59	15.71	9.32	7.07	4.58	8.12	12.42	62.46				
25	NT2RP2001943	14.71	11.45	18.53	11.8	23.77	7.53	6.86	10.79	7.88				
	NT2RP2001946	20.64	20.42	25.84	8.64	12.56	9.33	5.24	6.37	7.54	**	**	-	-
	NT2RP2001947	6.28	10.5	8.57	7.88	5.98	1.91	4.36	3.2	4.16	*		-	
	NT2RP2001956	21.67	10.17	24.79	16.61	12.12	14.12	16.82	19.29	13.33				
	NT2RP2001969	201.65	149.79	301.42	176.35	143.62	157.42	143.79	183.78	182.57				
	NT2RP2001976	4.33	7.95	17.7	7.78	4.49	4.49	2.81	5.46	10.35				
30	NT2RP2001978	14.35	12.6	20.91	12.02	5.57	5.03	3.1	3.56	4.91	**		-	
	NT2RP2001985	13.77	15.22	14.24	9.55	5.93	6.64	4.89	6.51	5.47	**	**	-	-
	NT2RP2001991	37.12	36.77	50.48	37.16	44.84	28.4	28.39	20.28	24.2	*		-	
	NT2RP2001997	13.99	12.07	14.92	11.27	12.08	8.47	13.49	12.5	12.19				
	NT2RP2002015	3.95	7.2	5.18	7.51	4.61	2.96	3.72	6.25	2.61				
	NT2RP2002017	14.92	13.42	18.23	7.3	5.04	8.42	3.4	6.85	8.43	**	*	-	-
35	NT2RP2002025	39.84	48.8	68.43	33.02	35.2	30.44	23.31	31.13	31.97				
	NT2RP2002030	28.92	28.65	40.93	26.27	23.9	23.74	29.55	36.7	30.22				
	NT2RP2002032	41.77	39.39	41.34	31.44	32.7	25.79	15.28	25.82	17.93	**	**	-	-
	NT2RP2002033	7.84	6.18	11.1	7.44	5.25	3.93	2.63	4.43	2.72	*		-	
	NT2RP2002041	63.15	38.11	58.94	48.43	54.38	37.99	33.68	26.24	30.79	*		-	
	NT2RP2002046	19.79	18.39	22.62	13.61	15.41	9.7	20.12	17.67	17.54	*		-	
40	NT2RP2002047	15.96	12.18	14.55	9.42	12.66	9.04	9.26	8.52	3.77	*		-	
	NT2RP2002050	6.47	7.96	8.94	7.07	2.69	4.33	1.22	3.6	2.62	**		-	
	NT2RP2002052	16.16	10.25	17.67	10.56	6.94	8.28	6.44	8.83	9.43				
	NT2RP2002058	10.7	8.12	13.82	9.12	8.72	4.7	3.71	8.14	3.23				
	NT2RP2002060	7.22	6.32	13.21	7.18	3.78	3.46	2.87	4.32	2.16				
	NT2RP2002063	27.7	18.33	28	16.06	12.42	6.01	14.89	17.06	15.58	*	*	-	-
45	NT2RP2002066	14.12	10.25	16.69	11.69	8.3	5.49	7.86	7.37	6.79	*		-	
	NT2RP2002070	47.82	69.53	26.7	25.88	50.9	22.08	11.62	12.95	13.2	*		-	
	NT2RP2002076	50.79	54.44	56.91	34.7	35.7	26.78	21.75	13.06	12.15	**	**	-	-
	NT2RP2002078	3.82	6.23	12.24	6.03	3.97	4.06	0.81	5.31	3.09				
	NT2RP2002079	13.68	8.15	17.98	9.69	6.98	8.2	5.55	8.29	6.41				
	NT2RP2002089	11.59	12.19	18.2	8.81	6.98	5.14	5.57	8.29	5.75	*	*	-	-
50	NT2RP2002105	15.09	12.54	15.38	10.34	7.26	6.01	5.5	9.12	6.68	*	**	-	-
	NT2RP2002115	15.39	14.23	22.91	11.04	12.24	9.95	11	13.28	11.06				
	NT2RP2002124	59.86	56.27	77.47	56.59	47.35	36.88	55.21	47.08	46.87				
	NT2RP2002137	8.42	8.75	8.28	6.56	6.55	3.48	2.47	3.09	3.68	*	**	-	-
	NT2RP2002139	93.62	131.25	91.96	52.77	52.45	52.95	81.88	52.45	48.22	*		-	
	NT2RP2002154	6.04	5.02	12.4	5.24	1.47	5.98	2.02	3.26	3.03				
55	NT2RP2002155	16.04	17.28	22.84	17	10.94	11.7	12.12	14.76	13.07				
	NT2RP2002172	26.05	32.14	34.35	17.74	14.37	15.16	12.78	22.56	16.28	**	*	-	-
	NT2RP2002185	13.73	14.4	21.2	14.18	14.86	10.72	12.32	12.52	11.2				

Table 425

	NT2RP2001347	539.84	514.01	628.74	565.74	422.96	416.2	495.32	585.14	429.89				
5	NT2RP2001366	38.71	33.77	43.04	29.42	28.24	19.52	18.19	26.65	21.17	*	*	-	-
	NT2RP2001378	8.65	6.18	12.14	10.82	6.77	5.51	3.49	4.36	4.96				
	NT2RP2001381	20.33	22.01	25.11	15	9.62	10.21	6.87	12.31	12.9	**	**	-	-
	NT2RP2001388	23.09	21.55	19.16	14.87	11.09	13.2	20	18.4	14.87	**		-	-
	NT2RP2001391	19.47	21.48	20.82	19.07	15.36	11.27	15.83	12.93	10.03	*		-	-
	NT2RP2001392	6.22	2.39	9.23	6.23	2.48	4.35	2.68	3.41	3.72				
10	NT2RP2001394	9	9.52	18.06	7.33	4.64	3.55	4.34	5.54	7.88				
	NT2RP2001397	5.52	4.95	15.76	7.71	3.52	1.79	2.61	4.45	3.74				
	NT2RP2001400	33.05	35.87	41.2	15.06	15.67	8.49	5.11	9.86	5.55	**	**	-	-
	NT2RP2001408	22.02	23.86	22.82	11.45	12.19	9.24	10.98	11.38	13.8	**	**	-	-
	NT2RP2001420	13.99	18.37	19.98	12.59	10.04	8.9	9.18	10.79	13.16	*	*	-	-
	NT2RP2001423	138.22	161.12	169.36	120.84	140.16	121.74	142.58	124.54	152.09				
15	NT2RP2001427	11.99	17.22	16.25	9.35	7.9	6.74	7.44	6.11	3.28	*	**	-	-
	NT2RP2001428	11.05	7.81	14.33	11.13	3.4	8	3.33	6.84	8.32				
	NT2RP2001436	28.98	21.86	34.92	22.78	25.35	19.1	15.23	27.9	27.44				
	NT2RP2001440	21.88	12.74	27.62	14.71	9.47	17.13	15.85	24.15	14.86				
	NT2RP2001445	9	10.8	19.43	10.91	10.54	6.62	8.66	8.73	7.21				
	NT2RP2001449	28.07	34.09	40.82	25.02	19.29	16.65	14.58	19.57	13.99	*	*	-	-
20	NT2RP2001450	20.8	28.19	27.38	18.17	18.42	8.85	15.67	22.32	18.66				
	NT2RP2001467	77.4	117.05	74.08	41.14	84.21	47.44	46.82	38.8	29.18	*		-	-
	NT2RP2001489	12.38	11.13	18.21	9.61	8.91	7.86	5.16	5.66	4.91	*		-	-
	NT2RP2001480	16.58	12.17	23.4	14.68	6.35	9.1	6.53	11.37	10.31				
	NT2RP2001495	5.39	5.05	10.71	5.62	3.39	3.05	2.02	2.99	3.13				
	NT2RP2001499	31.27	46.77	45.33	29.35	33.83	17.6	16.09	21.38	15.2	*		-	-
25	NT2RP2001506	7.58	7.79	10.57	9.25	7.24	4.11	3.34	2.96	2.75	**		-	-
	NT2RP2001508	10.21	8.66	19.01	9.32	5.78	3.48	10.19	10.43	10.1				
	NT2RP2001511	18.18	10.36	14.04	8.06	6.96	5.65	4.43	6.31	7.52	*	*	-	-
	NT2RP2001514	8.2	7.15	8.01	6.7	3.51	2.31	2.13	3.46	3.84	**		-	-
	NT2RP2001520	8.17	6.6	9.18	8.28	6.18	3.08	5.09	4.32	4.95	*		-	-
	NT2RP2001526	15.63	17.52	16.82	11.75	3.98	10.59	5.68	10.57	11.64	*	*	-	-
30	NT2RP2001529	18.96	9.11	22.28	13.03	8.09	10.02	6.2	8.22	8.57				
	NT2RP2001536	14.16	10.36	18.6	12.88	11.79	8.28	7.74	11.37	11.72				
	NT2RP2001538	8.11	6.47	12.78	7.67	5.34	3.77	4	4.36	2.56	*		-	-
	NT2RP2001547	9.99	10.57	12.92	9.74	7.58	3.49	4.05	3.74	4.33	**		-	-
	NT2RP2001560	11.5	10.74	9.02	6.82	5.56	7.18	6.87	5.66	4.21	*	*	-	-
	NT2RP2001562	17.39	15.7	17.62	10.84	8.02	7.21	8.92	9.2	10.24	**	**	-	-
35	NT2RP2001566	14.26	13.81	10.49	11.53	7.07	4.68	7.24	4.21	4.54	**		-	-
	NT2RP2001569	25.02	15.42	32.18	22.03	11.09	26.81	24.28	25.39	21.94				
	NT2RP2001576	129.89	178.91	114.02	47.3	49.96	55.77	33.67	55.78	72.18	*	*	-	-
	NT2RP2001581	20.4	22.7	22.23	13.8	17.88	10.54	10.57	16.71	15	*	*	-	-
	NT2RP2001597	17.49	9.3	13	11.25	8.35	11.5	4.54	4.57	3.83	*		-	-
	NT2RP2001601	376.46	416.73	382.11	192.76	196.04	131.77	252.12	219.45	367.42	**		-	-
40	NT2RP2001613	6.29	5.75	7.26	4.54	4.25	3.06	3.72	3.99	3.96	*	**	-	-
	NT2RP2001628	20.46	20.27	25.19	14.18	13.5	11.94	11.88	13.91	12.32	**	**	-	-
	NT2RP2001634	4.72	7.04	6	7.61	3.83	2.93	4.15	1.66	3.09	*		-	-
	NT2RP2001635	20.18	16.77	22.18	15.58	7.6	10.65	10.91	15.44	14.53	*	*	-	-
	NT2RP2001660	27.92	47.45	35.98	27.8	15.26	22.85	16.57	20.69	20.06	*		-	-
	NT2RP2001662	141.98	137.52	151.45	151.4	116.43	136.78	154.6	188.28	148.8				
45	NT2RP2001663	102.28	119.36	140.87	48.03	51.07	39.15	80.45	80.81	83.65	**	*	-	-
	NT2RP2001672	6.83	9.61	11.14	6.88	4.73	5.02	7.16	6.7	5.29				
	NT2RP2001675	7.51	11.75	9.95	6.41	3.17	2.8	4.39	4.7	4.26	*	*	-	-
	NT2RP2001677	18.83	22.14	19.19	10.64	6.34	6.91	9.72	6.83	7.68	**	**	-	-
	NT2RP2001678	10.06	8.87	8.82	7.97	6.41	4.67	9.59	5.34	9.32	*		-	-
	NT2RP2001683	88.93	103.45	119.49	195.62	102.44	134.62	102.34	95.16	65.68				
50	NT2RP2001699	36.15	36.2	44.96	18.8	7.95	13.41	6.7	9.44	13.11	**	**	-	-
	NT2RP2001707	6.77	3.9	7.75	5.48	4.72	5.02	3.95	4.21	3.45				
	NT2RP2001720	8.16	6.41	9.93	7.22	8.68	5.49	4.59	6.11	5.89				
	NT2RP2001721	33.41	34.55	34.5	25.33	18.3	17.93	32.85	38.18	33.9	**		-	-
	NT2RP2001740	64.22	76.89	65.2	34.74	42.49	31.73	26.06	24.24	41.05	**	**	-	-
	NT2RP2001748	6.3	15.07	7.65	6.61	1.99	2.37	4.7	3.94	2.45				
55	NT2RP2001766	8.56	8.68	5.59	6.1	3.13	5.08	4.74	3.64	4.01	*		-	-
	NT2RP2001762	4.42	4.43	4.61	6.06	1.37	4.5	3.19	3.5	2.83	**		-	-
	NT2RP2001768	7.43	5.23	6.6	7.86	2.49	4.84	3.03	5.68	3.45				

Table 424

	NT2RP2000863	9.23	2.37	9.61	12.6	5.57	10.29	2.95	6.5	2.03			
	NT2RP2000880	6.89	4.15	10.09	8.75	3.55	3.61	3.32	7.64	2.92			
5	NT2RP2000892	6.48	7.22	10.47	-6.68	3.59	2.44	2.63	2.13	2.32	**	-	
	NT2RP2000894	6.84	8.43	12.75	7.7	5.61	3.86	6.32	3.41	5.03			
	NT2RP2000903	14.72	12.47	10.22	6.37	7.06	6.25	6.3	4.12	3.45	*	**	-
	NT2RP2000906	7.83	5.78	8.49	5.67	3.5	4.08	5.51	1.9	1.89	*	*	-
	NT2RP2000910	7.08	7.41	7.6	9.95	3.31	4.88	6.74	6.36	7.22			
10	NT2RP2000931	17.23	16.66	19.14	9.83	11.03	7.29	5.66	13.37	8.87	**	*	-
	NT2RP2000932	4.85	5.4	12.51	5.58	2.98	2.75	2.51	3.48	1.84			
	NT2RP2000938	32.12	38.62	41.75	20.34	22.26	19.86	28.05	24.64	29.99	**	*	-
	NT2RP2000943	6.65	4.28	8.98	6.63	2.06	1.49	2.39	2.3	2.44	*		-
	NT2RP2000957	8.82	7.9	11.5	5.58	3.81	3.6	2.18	2.83	1.56	*	**	-
	NT2RP2000958	5.64	6.03	15.9	6.17	2.43	2.74	1.92	2.26	3.56			
15	NT2RP2000959	11.75	9.04	10.68	6.59	3.84	2.64	4.69	3.69	5.66	*	**	-
	NT2RP2000965	4.05	2.66	2.8	5.91	0.94	1.41	2.34	1.56	2.95			
	NT2RP2000970	58.85	65.2	74.74	64.15	40.74	92.64	72.54	97.39	89.84			
	NT2RP2000973	5.19	3.92	9.39	8.06	5.68	3.54	3.01	9.76	7.5			
	NT2RP2000985	3.85	2.51	4.33	4.79	3.19	4.55	1.82	2.45	2.12			
20	NT2RP2000987	39.38	36.27	32.68	33.06	40.49	26.38	33.61	26.04	27.01			
	NT2RP2000997	9.33	8.58	9.52	9.72	3.47	7.24	9.63	9.44	11.63			
	NT2RP2001024	12.74	7.92	18.67	6.22	2.48	2.6	4.96	3.99	3.75	*	*	-
	NT2RP2001028	38.71	37.69	41.88	33.98	15.8	30.64	34.47	25.69	33.62			
	NT2RP2001036	10.56	11.08	9.13	9.7	2.35	6.59	6.06	3.75	3.17	**	-	
	NT2RP2001039	29.77	24.74	30.19	17.85	11.75	29.98	20.42	28.95	23.24			
	NT2RP2001044	111.16	103.9	116.83	65.77	42.98	31.59	30.2	74.24	42.92	**	*	-
25	NT2RP2001056	16.47	11.38	12.03	9.34	5.04	8.68	12.51	8.41	8.95			
	NT2RP2001065	12.87	18.54	21.09	9.6	6.83	8.58	6.23	6.54	8.28	*	*	-
	NT2RP2001067	18.27	13.06	19.79	9.25	5.26	3.58	-5.5	6.23	8.66	*	*	-
	NT2RP2001070	4.55	6.03	7.25	6.14	2.9	2.2	3.37	2.83	1.95	*		-
	NT2RP2001081	11.96	15.94	13.97	8.19	3.81	5.25	5.58	4.15	6.83	**	**	-
	NT2RP2001087	65.9	61.9	76.08	39.7	16.73	50.09	38.08	21.24	22.36	*	**	-
30	NT2RP2001094	12.2	8.99	11.02	14.61	10.72	9.52	11.02	11.03	11.05			
	NT2RP2001119	4.16	4.58	7.07	10.85	1.99	7.07	4.12	3.63	2.63			
	NT2RP2001127	11.36	13.12	3.9	16.08	4.34	7.73	5.38	7.85	3.42			
	NT2RP2001133	5.67	4.14	5.86	10.97	6.13	8.56	8.68	4.41	3.17			
	NT2RP2001137	49.8	41.1	45.81	39.26	21.06	24.92	25.21	45.63	17.6	*	-	
	NT2RP2001142	5.27	5.53	5.06	5.3	2.26	3.88	2.41	3.56	3.78	*	-	
35	NT2RP2001149	6.92	5.61	5.45	13.36	3.82	6.27	6.19	4.78	4.91			
	NT2RP2001168	6.12	7.49	5.66	8.34	5.85	5.09	5.44	4.63	5.1			
	NT2RP2001173	4.74	3.23	8.28	7.11	2.4	3.67	4.61	6.54	6.37			
	NT2RP2001174	44.63	21.39	37.72	44.99	18.88	46.31	52.67	53.94	56.67	*	+	
	NT2RP2001184	7.2	4.99	10.33	8.93	4.09	3.96	4.88	4.86	5.6			
40	NT2RP2001196	188.27	171.63	278.21	277.29	156.05	217.9	310.45	339.15	334.54	*	+	
	NT2RP2001200	5.84	6.42	11.03	6.08	6.9	5.52	4.06	2.49	4.45			
	NT2RP2001218	12.41	10.49	14.66	10.34	6.96	7.23	9.11	8.8	8.23	*	-	
	NT2RP2001223	10.09	12.26	12.04	13.71	9	9.77	10.94	7.86	9.72			
	NT2RP2001226	295.34	151.71	253.15	264.52	155.79	205.15	225.76	302.68	279.91			
	NT2RP2001227	9.29	9.83	15.24	10.83	5.91	8.02	6.71	6.54	8.43			
	NT2RP2001232	6.25	9.05	13.72	6.11	9.52	4.37	3.08	5.35	5.73			
45	NT2RP2001233	17.2	11.64	26.2	12.06	8.02	7.48	8.66	8.69	6.62			
	NT2RP2001245	42.01	30.76	56.72	22.72	58.34	32.96	29.15	30.96	32.88			
	NT2RP2001246	5.12	5.92	8.66	6.91	4.33	3.68	3.64	7.01	3.2			
	NT2RP2001268	76.86	63.92	101.42	88.4	109.5	90.32	72.85	73.19	77.59			
	NT2RP2001270	10.64	12.74	14.24	12.74	10.24	11.95	10.68	9.82	10.94			
	NT2RP2001276	23.31	16.76	30.97	21.15	16.68	10.08	14.57	16.7	14.66			
50	NT2RP2001277	24.91	10.73	18.18	16.47	6.03	6.24	22.15	25.09	49.12			
	NT2RP2001290	8.95	7.82	10.48	8.48	7.41	5.73	5.82	9.63	10.81			
	NT2RP2001295	14.82	7.48	15.13	8.28	5.62	4.48	5.33	6.4	4.82			
	NT2RP2001297	8.15	6.67	10.81	9.36	7.32	4.92	4.17	4.98	6			
	NT2RP2001301	47.34	44.45	63.71	42.51	38.25	33.89	33.81	29.8	26.52	*	-	
	NT2RP2001312	8.43	10.98	14.02	7.43	5.35	5.08	6.81	4.46	4.32	*	*	-
55	NT2RP2001327	20.88	28.57	34.54	16.53	15.03	15.84	11.1	20.64	9.65	*	-	
	NT2RP2001328	10.32	3.77	9.25	9.2	3.81	6.74	4.44	5.56	7.15			
	NT2RP2001341	3499.3	2809.3	4099.1	1364.9	1644.2	898.17	1201.7	2570.4	2392	**	-	

Table 423

	NT2RP2000448	25.71	13.78	32.06	17.1	19.22	17.26	5.88	10.24	7.42	*	-
5	NT2RP2000459	9.24	9.95	9.52	9.52	4.14	4.36	2.59	6.53	3.6	*	-
	NT2RP2000479	6.63	5.55	7.74	7.89	3.36	5.52	5.8	5.59	6.42	*	-
	NT2RP2000498	11.27	6.84	9.95	5.98	2.48	3.08	4.73	2.88	2.16	*	-
	NT2RP2000503	14.72	13.18	15.3	32.52	21.1	19.93	18.56	11.89	14.4	*	-
	NT2RP2000510	7.33	5.8	8.13	5.52	2.28	3.46	4.2	5.04	2.87	*	-
	NT2RP2000514	4.57	2.69	2.25	7.44	1.96	3.28	5.71	4	3.87	*	-
10	NT2RP2000516	6.54	4.62	8.39	7.51	4.34	6.37	3.47	4.35	4.95	*	-
	NT2RP2000523	4.11	2.38	11.01	7.29	1.39	3.22	3.15	2.27	3.38	*	-
	NT2RP2000533	2.68	1.37	7.63	4.63	1.22	2.09	0.79	2.38	0.48	*	-
	NT2RP2000540	6.16	5.88	8.08	5.48	4.38	3.8	3.01	4.98	1.52	*	-
	NT2RP2000547	7.42	7.44	10.91	6.02	3.13	4.89	2.98	4.52	3.39	*	-
	NT2RP2000557	49.31	60.55	82.94	43.46	53.17	74.9	83.43	53.58	63.67	*	-
15	NT2RP2000558	18.68	19.95	27.42	20.69	14.13	19.17	24.64	17.82	18.81	*	-
	NT2RP2000564	0.43	0.11	2.04	5.89	0.93	1.24	2.05	1.59	0.53	*	-
	NT2RP2000565	13.78	7.23	14.74	12.39	9.94	11.42	3.9	7.3	8.83	*	-
	NT2RP2000583	10.87	6.32	13.24	7.55	3.8	5.68	4.41	5.83	9.68	*	-
	NT2RP2000591	7.05	5.42	12.75	6.46	3.08	3.75	3.89	6.69	2.85	*	-
	NT2RP2000599	45.38	33.04	53.33	45.68	38.74	39.04	49.46	50.96	42.92	*	-
20	NT2RP2000601	5.8	2.35	8.26	6.01	2.72	4.74	3.53	6.5	4.71	*	-
	NT2RP2000603	10.6	10.97	14.76	12.57	11.91	10.27	11.35	11.32	10.57	*	-
	NT2RP2000610	16.79	27.41	21.69	12.86	12.08	9.62	7.92	6.77	7.19	*	-
	NT2RP2000614	22.5	27.77	21.5	15.84	12.38	13.2	11.46	7.34	10.15	*	-
	NT2RP2000616	15.9	10.73	17.88	14.04	10.65	8.26	8.53	13.65	30.31	*	-
	NT2RP2000617	7	4.89	10.44	5.79	2.49	3.27	2.33	3.91	6.08	*	-
25	NT2RP2000623	30.93	27.18	42.39	23.25	19.11	22.34	22.03	24.06	17.67	*	-
	NT2RP2000634	5.41	5.13	8.98	7.02	5.17	5.22	4.91	6.31	5.96	*	-
	NT2RP2000636	6.22	7.05	7.96	5.06	2.33	3.65	3.42	4.64	2.62	*	-
	NT2RP2000638	5.4	5.87	8.77	7.49	5.36	5.46	5.01	4.21	4.57	*	-
	NT2RP2000644	6.06	11.57	13.6	11.21	5.35	7.7	5.1	5.24	4.14	*	-
	NT2RP2000649	69.57	64.76	72	79.36	45.65	56.22	93.91	57.37	68.85	*	-
30	NT2RP2000652	14.9	10.94	49.84	10.88	5.91	11.87	6.2	5.26	32.14	*	-
	NT2RP2000656	12.57	7.15	13.54	6.29	4.4	6.3	1.69	9.03	14.78	*	-
	NT2RP2000658	5.45	4.52	10.98	5.11	2.13	3.91	1.81	2.12	1.71	*	-
	NT2RP2000663	4.6	4.28	7.43	5.58	2.25	5.97	4.47	2.44	1.11	*	-
	NT2RP2000664	10.9	7.61	12.36	9.42	3.88	8.64	13.44	10.25	9.81	*	-
	NT2RP2000668	33.38	33.11	48.17	36.56	39	37.32	34.56	25.63	30.2	*	-
35	NT2RP2000678	6.12	9.03	9.25	9.6	7.57	6.65	6.88	6.02	5.34	*	-
	NT2RP2000694	16.87	11.51	13.29	13.35	8.47	12.93	22.9	14.8	18.71	*	-
	NT2RP2000704	2.84	3.9	14.85	6.19	2.87	7.29	2.98	6.45	4.78	*	-
	NT2RP2000710	8.84	8.7	19.22	7.25	3.74	6.4	5.24	6.81	6.9	*	-
	NT2RP2000712	5.84	5.25	16.26	7.12	3.27	5.34	2.82	4.15	2.23	*	-
	NT2RP2000715	23.49	31.67	27.44	16.78	13.7	10.05	7.2	11.02	9.66	*	-
40	NT2RP2000720	14.6	11.95	17.99	9.67	7.04	6.8	9.67	8.11	6.87	*	-
	NT2RP2000731	130.46	114.16	171.06	139.58	136.44	144.08	91.22	91.1	91.43	*	-
	NT2RP2000739	10.56	11	13.93	9.61	7.28	5.99	9.08	6.03	5.15	*	-
	NT2RP2000748	12.42	9.17	9.8	13.86	6.05	5.01	6.86	4.28	5.36	*	-
	NT2RP2000749	11.23	7.84	14.76	7.12	4.44	6.62	1.43	4.83	4.58	*	-
	NT2RP2000758	18.71	12.4	22.27	12.06	7.43	9.25	3.34	5.27	4.96	*	-
45	NT2RP2000764	8.83	9.65	15.17	9.68	4.51	6.36	4.78	3.89	4.68	*	-
	NT2RP2000766	39.21	45.57	54.49	39.05	26.43	20.35	16.67	26.46	19.71	**	-
	NT2RP2000777	37.43	31.1	43.71	20.28	16.78	21.16	3.56	8.61	6.24	**	-
	NT2RP2000786	10.79	9.6	16.45	9.08	5.82	3.61	8.49	11.1	8.19	*	-
	NT2RP2000793	8.96	7.43	14.72	5.59	4.46	3.27	3.41	3.35	2.39	*	-
	NT2RP2000796	13.41	11.17	11.19	12.52	8.7	5.88	4.89	4.55	5.82	**	-
50	NT2RP2000809	9.51	8.66	16.69	9.15	5.7	10.07	8.67	9.07	9.06	*	-
	NT2RP2000812	10.78	5.64	19.76	9.49	6.33	7.02	9.58	10.65	9.48	*	-
	NT2RP2000814	2.73	2.98	11.46	6.79	1.46	2.03	1.53	1.11	1.73	*	-
	NT2RP2000816	5.26	5.27	12.58	5.83	4.32	3.54	2.61	2.6	1.81	*	-
	NT2RP2000818	64.67	52.94	88.69	38.83	38.29	31.82	36.98	34.96	36.14	**	-
	NT2RP2000819	10.64	9.84	11.3	8.11	4.61	3.58	6.06	2.23	3.74	*	-
55	NT2RP2000841	19.27	13.42	17.68	15.7	9.96	11	12.88	9.6	13.79	*	-
	NT2RP2000842	7.9	5.25	9.24	8.2	4.88	2.43	6.17	3.9	4.72	*	-
	NT2RP2000845	1.64	3.58	10.06	6.26	3.24	5.16	2.58	3.32	3.23	*	-

Table 422

	NT2RP2000108	10.87	8.33	17.31	10.58	6.71	6.74	4.11	5.62	13.35			
5	NT2RP2000114	14.61	10.83	24.65	15.49	7.8	10.66	10.74	13.91	10.18			
	NT2RP2000116	12.92	13.17	18.63	12.07	8.1	10.87	4.57	5.57	3.24	**	**	-
	NT2RP2000119	50.32	47.2	56.58	31.02	20.71	31.82	30.15	29.31	28.57	**	**	-
	NT2RP2000120	7.9	9.18	10.99	6.62	6.21	6.5	4.81	5.84	4.99	*	*	-
	NT2RP2000126	27.18	24.32	32.55	18.42	11.34	17.52	24.37	18.13	19.51	*		-
	NT2RP2000133	6.31	5.61	8.16	7.63	3.91	7.78	7.66	10.7	6.35			
10	NT2RP2000147	12.98	10.72	23.8	11.64	4.41	11.41	3.62	6.37	10.62			
	NT2RP2000153	4.49	6.28	13.18	7.15	3.26	3.2	2.66	3.8	4.92			
	NT2RP2000156	7.61	6.37	12.56	6.43	4.23	4.03	2.05	6.21	1.59			
	NT2RP2000157	64.87	55.86	69.1	57.96	47.99	49.02	51.12	61.28	54.87			
	NT2RP2000161	8.86	7.44	9.1	8.5	4.93	4.27	4.28	4.21	2.7	**		-
	NT2RP2000168	8.64	8.81	11.56	8.06	4.98	11.13	4.05	3.85	9.74			
15	NT2RP2000173	5.87	12.11	13.07	7.31	10.56	7.74	13.29	16.36	4.1			
	NT2RP2000175	6.28	5.9	5.57	9.33	4.37	3.63	5.8	5.65	4.1			
	NT2RP2000178	7.4	9.25	18.07	7.99	4.34	5.36	2.6	5.71	4.5			
	NT2RP2000183	5.74	7.89	15.69	7.74	4.42	5.84	3.67	6.86	4.04			
	NT2RP2000195	16.12	12.54	17.86	16.63	13.32	13.8	5.12	8.05	4.54	**		-
	NT2RP2000204	22.18	23.46	35.46	16.84	14.7	7.3	7.36	8.83	4.75	*		-
20	NT2RP2000205	306.16	263.17	279.79	153.75	186.84	122.03	44.13	69.43	57.15	**	**	-
	NT2RP2000208	5.24	2.8	6.75	5.38	2.18	0.7	2.69	2.06	2.36			
	NT2RP2000224	24.97	24.03	27.21	17.41	19.54	13.73	10.21	10.44	6.58	*	**	-
	NT2RP2000230	26.52	31.48	20.02	29.78	28.26	14.04	27.93	14.94	32.18			
	NT2RP2000231	102.91	95.51	132.32	51.89	45.81	56.22	28.1	59.69	105.21	**		-
	NT2RP2000232	18.39	14.23	28.59	15.4	9.71	12.46	6.13	9.73	7.72	*		-
25	NT2RP2000233	7.64	7.96	15.89	7.58	4.07	3.39	2.14	3	2.26	*		-
	NT2RP2000239	66.01	139	172.9	85.94	84.53	44.49	55.89	55.75	25.78			
	NT2RP2000240	5.47	3.94	10.61	4.6	1.25	2.17	1.78	2.32	0.43			
	NT2RP2000248	6.07	5.52	8.36	6.23	4.28	2.64	2.65	2.47	2.64	**		-
	NT2RP2000256	16.57	14.62	18.18	13.37	8.16	8.91	7.89	5.94	4.14	*	**	-
	NT2RP2000257	39.73	47.22	36.2	40.72	25.44	29.27	36.28	29.33	25.56			
30	NT2RP2000258	13.09	11.66	21.23	12.78	8.27	8.08	5.57	13.01	9.17			
	NT2RP2000261	8.27	7.35	17.74	7.68	2.93	4.17	3.42	5.31	3.64			
	NT2RP2000270	3.96	6.69	13.81	17.92	6.26	3.85	3.35	9.95	4.71			
	NT2RP2000274	91.36	80.94	71.7	59.79	55.47	43.9	66.54	63.63	61.61	*	*	-
	NT2RP2000277	4.35	3.88	5.68	6.38	1.67	1.79	1.78	1.55	2.23	**		-
	NT2RP2000279	7.44	6.18	9.49	6.39	2.22	2.73	3.81	2.76	3.21	*		-
35	NT2RP2000283	5.12	8.14	7.43	6.14	1.91	2.31	3.36	2.12	4.13	*		-
	NT2RP2000284	10.29	7.99	9.75	9.32	5.86	5.55	9.02	7.12	9.38			
	NT2RP2000289	21.4	17.79	32.55	15.37	7.32	16.77	22.06	23.42	27.43			
	NT2RP2000297	4.51	6.32	10.63	6.94	2.23	3.64	3.53	4.19	2.55			
	NT2RP2000298	8.05	8.79	13.64	5.87	3.8	2.92	2.96	3.04	1.11	*	*	-
	NT2RP2000310	80.52	91.37	87.72	32.16	27.21	29.67	20.91	14.48	20.59	**	**	-
40	NT2RP2000327	9.08	9.43	13.33	5.88	3.81	4.9	2.31	2.04	2.13	*	**	-
	NT2RP2000328	21.82	17.34	15.94	8.37	8.84	6.89	5.86	4.83	2.63	**	**	-
	NT2RP2000329	17.76	16.87	20.08	10.58	7.07	6.38	10.73	6.62	8.14	**	**	-
	NT2RP2000333	8.83	7.98	9.31	8.74	1.76	4.67	8.04	7.67	6.48			
	NT2RP2000337	26.86	26.84	31.77	17.74	6.47	12.94	6.6	13.35	15.87	*	**	-
	NT2RP2000346	6.79	9.29	9.75	14.24	7.42	4.48	6.69	17.82	10.4			
45	NT2RP2000357	14.11	13.18	22.49	5.83	2.27	3.95	3.56	4.39	2.66	*	*	-
	NT2RP2000358	9.23	5.13	8.16	5.34	2.02	3.61	4.07	3.18	2.94	*		-
	NT2RP2000366	5.06	4.81	6.09	5.12	1.19	2.67	2.27	2.31	1.55	**		-
	NT2RP2000369	29.84	26.23	39.9	17.03	12.11	15.49	15.78	7.96	15.06	*	*	-
	NT2RP2000376	26.15	35.86	51.75	24.31	15.42	18.34	35.37	27.75	35.03			
	NT2RP2000384	11.62	6.66	9.45	10.5	5.28	8.79	9.06	9.09	12.53			
50	NT2RP2000396	2.03	7.18	10.2	4.43	3.16	1.53	2.05	5.24	2.77			
	NT2RP2000412	13.13	15.96	18.35	14.6	6.73	8.07	6.73	16.17	12.59			
	NT2RP2000414	19.9	19.23	18.62	8.95	6.33	5.66	6.29	8.15	6.52	**	**	-
	NT2RP2000420	5.15	7.34	11.45	10.83	6.03	5.33	5.71	4.93	7.76			
	NT2RP2000422	10.72	8.26	11.6	8.9	4.28	5.31	4.96	4.79	6.17	*		-
	NT2RP2000426	7.55	6.99	6.8	4.75	1.02	1.72	2.91	2.26	1.94	*	**	-
55	NT2RP2000428	8.74	4.22	5.73	4.39	1.17	2.36	8.21	2.73	3.16			
	NT2RP2000438	12.17	11.38	11.77	10.69	3.56	5.65	4.51	2.36	2.93	**		-
	NT2RP2000447	12.75	6.94	10.08	6.36	4.33	7.41	3.21	7.55	5.61			

Table 421

	NT2RP1001302	13.32	15.35	23.89	11.51	7.14	10.86	5.3	9.24	8.54	*	-
5	NT2RP1001310	78.58	77.83	107.95	40.93	36.67	35.03	13.95	16.52	15.88	**	-
	NT2RP1001311	12.5	10.19	20.17	13.97	8.87	11.71	8.61	11.48	9.2	*	-
	NT2RP1001313	47.33	35.38	51.8	18.55	17.75	13.75	4.5	6.14	3.62	**	-
	NT2RP1001324	13.11	11.52	14.26	8.91	7.65	4.65	7	7.07	7.14	*	-
	NT2RP1001349	8.69	8.34	8.01	6.09	3.96	4.38	3.75	1.87	3.49	**	-
	NT2RP1001361	9.54	13.7	11.81	11.62	6.33	6.68	6.92	4.67	4.5	*	-
10	NT2RP1001379	13.14	11.82	15.33	13.1	6.47	9.99	7.99	14.61	8.34	*	-
	NT2RP1001385	24.66	22.66	28.03	8.27	7.17	9.45	13.37	13.91	22.49	**	-
	NT2RP1001395	5.65	15.28	10.96	10.51	4.01	1.7	3.74	3.38	2.71	*	-
	NT2RP1001410	13.37	15.22	15.93	8.65	10.09	5.26	5.57	5.86	7.15	*	-
	NT2RP1001424	28.73	25.67	34.05	32.06	24.11	29.55	22.85	15.99	24.49	*	-
	NT2RP1001432	9.64	7.96	12.22	8.86	4.28	6.46	5.04	7.62	6.96	*	-
15	NT2RP1001449	19.34	21.85	20.51	10.44	6.36	4.06	9.88	5.66	6.23	**	-
	NT2RP1001457	23.3	38.81	37.3	24.2	13.72	10.77	10.68	7.09	13.11	*	-
	NT2RP1001459	13.65	11.76	16.02	12.39	4.68	8.62	6.06	9.05	6.34	*	-
	NT2RP1001466	8.39	4.23	8.84	7.44	4.36	4.01	3.12	6.63	4.14	*	-
	NT2RP1001475	11.89	17.85	12.86	11.05	10.38	6.52	5.05	12.24	6.54	*	-
	NT2RP1001482	15.19	20.93	20.15	19.85	11.71	18.03	11.8	14.09	12.75	*	-
20	NT2RP1001494	3.95	5.63	10.18	5.63	2.65	3.16	2.73	3.52	2.73	*	-
	NT2RP1001500	9.08	4.79	11.52	5.37	1.85	1.77	2.31	4.4	3.24	*	-
	NT2RP1001517	18.13	18.45	18.1	8.4	4.89	3.69	7.15	5.46	6.44	**	-
	NT2RP1001540	28.09	31.99	28.56	25.07	15.6	22.78	25.04	22.39	21.45	*	-
	NT2RP1001543	20.19	16.33	24.14	16.6	10.69	13.4	7.07	16.07	9.64	*	-
	NT2RP1001546	12.99	15.98	15.41	9.88	3.03	7.01	6.04	9.72	7.53	*	-
25	NT2RP1001550	8.32	14.7	14.31	7.95	15.82	5.57	6.4	9.76	9.43	*	-
	NT2RP1001553	5.3	7.23	13.14	7.37	4.11	4.11	3.68	3.77	3.88	*	-
	NT2RP1001555	54.76	52.6	55.46	31.87	21.44	34.31	43.17	41.32	32.44	**	-
	NT2RP1001563	39.08	33.85	48.64	38.88	21.62	23.18	31.1	28.26	33.8	*	-
	NT2RP1001589	3.96	1.69	1.05	5.34	1.73	1.64	1.66	1.23	2	*	-
30	NT2RP1001584	172.68	220.01	198.04	20.64	11.47	19.54	81.72	38.4	33.43	**	-
	NT2RP1001599	4.12	3.82	5.18	7.25	3.83	4.97	3.61	3.84	3.39	*	-
	NT2RP1001616	4.03	1.77	2.09	6.09	1.64	4	2.04	2.76	1.59	*	-
	NT2RP1001654	14.62	9.64	12.33	11.97	7.7	10.52	5.54	7.93	4.12	*	-
	NT2RP1001665	5.76	6.54	8.08	8.07	4.13	4.72	3.98	9.32	5.29	*	-
	NT2RP1001679	8.69	7.12	7.86	9.21	3.08	4.08	4.04	6.22	5.1	*	-
35	NT2RP1001681	18.99	18.82	24.49	14.39	5.58	9.22	11.12	13.68	12.99	*	-
	NT2RP1001694	6.52	3.33	5.96	7.17	2.61	3.23	4.87	4.47	3.87	*	-
	NT2RP2000001	6.24	4.74	8.49	6.13	5.97	4.89	6.42	5.79	3.58	*	-
	NT2RP2000006	9.03	5.67	13.4	11.28	7.14	5.76	7.62	8.34	6.39	*	-
	NT2RP2000007	10.8	6.31	17.13	11.11	4.89	9.86	6.74	8.3	8.67	*	-
	NT2RP2000008	5.65	3.68	9.46	7.12	2.84	3.88	3.74	5.6	5.37	*	-
	NT2RP2000010	6.84	5.53	11.05	6.98	2.63	4.35	3.2	3.54	2.21	*	-
40	NT2RP2000011	9.36	13.46	17.26	10.79	6.33	9.28	5.31	3.23	5.5	*	-
	NT2RP2000027	3.47	3.05	7.32	6.1	1.45	4.19	3.16	3	5.68	*	-
	NT2RP2000028	4.74	6.32	6.74	8.06	2.13	4.67	3.21	1.66	4.03	*	-
	NT2RP2000032	3.24	7.28	8.03	10.28	5.5	7.28	3.26	3.13	4.05	*	-
	NT2RP2000040	15.29	8.48	28.46	11.5	6.66	9.38	4.88	6.35	7.25	*	-
45	NT2RP2000042	21.14	15.46	35.04	11.42	8.01	11.22	12.48	11.99	17.82	*	-
	NT2RP2000045	5.83	3.94	13.32	7.43	3.18	3.46	3.62	5.2	3.95	*	-
	NT2RP2000051	13.61	9.89	23.1	13.04	8.32	9.72	10.33	13.1	11.12	*	-
	NT2RP2000054	16.51	12.89	19.57	11.63	9.66	11.02	9.64	7.65	7.91	*	-
	NT2RP2000056	6.15	6.7	14.22	6.37	4.1	4.09	7.75	7.98	8.74	*	-
	NT2RP2000057	7.7	5.49	9.31	7.63	5.41	5.64	12.9	3.34	4	*	-
	NT2RP2000067	7.64	15.12	15.54	18.12	16.74	10.35	9.45	6.54	9.35	*	-
50	NT2RP2000070	522.83	440.99	557.3	464.06	449.31	610.82	418.05	662.99	650.93	*	-
	NT2RP2000076	6.99	4.98	10.04	7.01	1.06	1.43	1.54	3.61	5.67	*	-
	NT2RP2000077	14.55	10.4	23.49	9.32	7.18	6.26	8.55	8.84	8.57	*	-
	NT2RP2000079	9.19	8.49	12.44	8.77	5.26	7.56	6.72	6.46	6.39	*	-
	NT2RP2000088	17.05	20.32	26.56	22.32	20.65	23.39	19.38	14.35	14.55	*	-
	NT2RP2000091	33.24	19.21	39.37	24.16	20.9	17.89	11.58	9.95	7.77	*	-
55	NT2RP2000092	0.22	2.2	2.32	4.37	1.08	1	2.25	1.85	1.49	*	-
	NT2RP2000097	10.19	11.77	12.54	11.39	5.79	8.17	7.35	3.3	4.01	**	-
	NT2RP2000098	14.34	13.24	18.96	13.26	6.07	9.3	5.46	9.38	18.63	*	-

Table 420

	NT2RP1000780	19.06	11.29	17.88	11.55	12.19	16.36	7.6	12.92	8.96			
	NT2RP1000782	3.8	1.98	4.82	5.26	2.11	4.2	3.73	3.14	3.38			
5	NT2RP1000796	23.21	12.88	16.72	13.54	7.34	8.03	3.16	8.05	3.72	*	-	
	NT2RP1000797	8.11	7.21	12.88	6.28	1.88	5.78	3.95	3.02	3.35	*	-	
	NT2RP1000800	6.36	7.42	8.45	6.35	6.28	4.56	5.14	4.53	6.09	*	-	
	NT2RP1000825	6.14	10.58	8.77	8.58	2.44	5.18	5.33	5.66	4.12			
	NT2RP1000833	6.32	4.62	5.45	7.5	3.18	3.17	4.57	14.3	3			
	NT2RP1000834	12.66	13.91	14.59	8.48	3.38	6.31	7.69	5.15	4.86	**	**	-
10	NT2RP1000836	6.87	3.86	10.29	7.94	2.62	4.87	5.98	4.8	8.86			
	NT2RP1000837	4.76	2.83	13.29	5.43	1.86	3.86	3.47	3.54	3.43			
	NT2RP1000846	14.34	10.47	22.36	10.56	7.41	9.61	13.44	9.9	11.55			
	NT2RP1000847	17.63	21.93	45.89	20.33	17.15	15.75	21.4	13.87	15.38			
	NT2RP1000851	3.61	3.71	7.21	5	2.22	2.83	3.04	2.47	2.36			
	NT2RP1000856	15.4	23.9	24.75	12.89	7.5	8.85	6.97	4.98	5.2	*	**	-
15	NT2RP1000860	5.61	4.82	11.39	9.47	4.26	7.06	6.83	5.9	5.93			
	NT2RP1000902	10.67	7.93	13.65	15.09	8.49	17.09	4.79	9.4	9.51			
	NT2RP1000903	14.49	11.64	18.42	14.24	5.67	14.38	12.23	14.45	21.38			
	NT2RP1000905	11.82	8.69	16.54	6.96	4.86	6.4	5.28	5.14	5.21	*	-	
	NT2RP1000915	63.69	60.39	85.78	55.83	55.49	63.74	77.2	80.56	84.89			
	NT2RP1000916	81.28	82.59	69.7	73.31	86.67	61.79	53.76	46.45	55.88	**	-	
20	NT2RP1000921	71.86	52.08	72.49	27.75	31.76	38.72	21.98	23.9	25.92	*	**	-
	NT2RP1000943	6.66	5.16	12.72	6.58	6.6	3.97	5.26	4.45	3.75			
	NT2RP1000944	36.56	43.79	53.16	29.61	24.97	30.98	38.1	30.4	32.9	*	-	
	NT2RP1000947	229.13	162.3	242.56	211.36	169.75	208.5	187.13	230.69	193.69			
	NT2RP1000954	8.73	6.8	11.31	9.14	2.72	5.18	2.45	4.94	9.97			
25	NT2RP1000958	12.83	9.08	19.99	13.3	9.58	12.08	6.34	10.95	18.7			
	NT2RP1000959	51.88	27.73	29.26	17.4	13.14	15.25	9.23	12.74	9.9	*	-	
	NT2RP1000966	30.03	20.71	21.11	23.97	20.83	19.34	9.71	14.32	8.82	*	-	
	NT2RP1000974	16.61	13.83	23.36	13.15	11.64	14.93	7.2	7.22	7.83	*	-	
	NT2RP1000980	85.21	82.27	107.69	76.1	98.09	99.97	80	57.54	67.02			
	NT2RP1000981	14.84	15.8	18.22	12.59	10.09	13.46	23.72	16.48	19.72	*	-	
30	NT2RP1000988	21.09	10.38	16.08	11.96	7.25	9.77	8.85	10.68	7.42			
	NT2RP1001002	23.42	12.11	26.95	15.71	9.49	11.37	4.76	9.9	15.94			
	NT2RP1001004	3.32	4.31	12.34	5.67	1.55	2.59	2.12	2.77	2.28			
	NT2RP1001007	5.36	4.39	9.57	6.91	2.92	2.64	2.01	3.93	1.56			
	NT2RP1001011	5.98	6.54	6.16	6.28	2.04	2.66	2.36	4.56	2.02	*	-	
	NT2RP1001013	168.04	170.77	232.56	168.44	147.11	154.85	176.26	171.79	185.19			
35	NT2RP1001014	13.02	16.54	16.26	8.7	8	7.08	6.95	5.62	4.04	**	**	-
	NT2RP1001020	12.63	15.72	12.22	15.74	11.78	16.3	13.8	9.3	14.27			
	NT2RP1001023	16.05	14.39	18.27	13.53	16.55	17.02	5.93	11.56	12.91			
	NT2RP1001027	5.55	8.55	17.95	9.25	6.73	6.98	2.42	5.11	5.43			
	NT2RP1001031	5.67	4.8	13.77	5.12	2.4	2.34	1.77	3.34	2.2			
	NT2RP1001033	7.93	6.3	11.72	6.43	4.31	2.18	2.83	5.21	2.62	*	-	
40	NT2RP1001042	63.59	58.82	65.69	59.83	59.66	45.94	71.71	83.48	74.44	*	+	
	NT2RP1001045	9.83	10.21	13.29	8.26	4.99	5.95	3.06	4.58	3.85	*	**	-
	NT2RP1001073	12.41	18.41	17.37	9.19	7.47	7.07	4.73	4.66	5.19	*	**	-
	NT2RP1001079	38.47	46.51	52.14	22.17	29.03	25.48	25.23	15.77	15.85	*	**	-
	NT2RP1001080	21.61	6.66	17.31	12.32	7.25	12.68	6.59	12.92	8.31			
	NT2RP1001113	57.87	45.81	82.08	49.33	42.8	51.52	16.27	50.38	44.76			
45	NT2RP1001159	7.51	4.75	14.77	8.2	4.64	7.14	4.3	5.24	3.99			
	NT2RP1001173	4.39	5.23	11.13	7.48	3.15	1.42	1.19	2.31	1.17			
	NT2RP1001176	52.66	45.96	57.66	32.33	25.13	21.71	19.56	17.05	21.12	**	**	-
	NT2RP1001177	10.79	9.96	12.22	5.4	2.87	3.22	4.03	4.39	5.3	**	**	-
	NT2RP1001185	14.18	11.86	11.53	9.31	7.59	5.91	7.44	8.55	7.87	*	**	-
	NT2RP1001199	13.52	13.24	12.28	9.53	6.55	5.34	4.8	3.41	3.43	*	**	-
50	NT2RP1001205	93.58	72.48	94.49	51.42	53.78	86.93	52.32	75.19	84.87			
	NT2RP1001215	17.01	15.57	25.94	10.31	11.47	12.47	9.75	16.75	14.31			
	NT2RP1001225	11.8	18.2	17.52	11.07	6.07	5.97	4.13	11.34	3.6	*	*	-
	NT2RP1001245	6.59	5.31	9.93	6.03	2.96	4.57	2.73	3.24	2.91	*	-	
	NT2RP1001247	6.32	3.32	10.27	4.35	3.87	3.01	3.42	3.7	1.09			
	NT2RP1001248	15.25	12.77	20.45	15.52	14.1	11.98	13.33	13.26	15.08			
55	NT2RP1001253	84.68	116.5	77.6	46.18	37.13	24.55	27.4	9.2	25.65	*	**	-
	NT2RP1001286	5.14	8.32	6.96	7.42	3.69	3.62	3.74	2.33	2.9	*	-	
	NT2RP1001294	57.97	32.23	60.31	62.38	43.54	78.24	53.21	40.65	43.49			

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	NT2RP1000386	37.77	48.73	51.62	24.07	21.15	29.99	24.61	14.84	15.66	*	**	-	-
	NT2RP1000407	20.7	11.19	17.51	22.94	11.29	15.13	8.7	15.9	29.09				
5	NT2RP1000409	24.93	20.8	32.63	23.46	10.36	22.79	11.36	15.76	11.35	*		-	
	NT2RP1000413	24.89	20.12	32.27	23.23	14.81	20.34	6.71	17.88	15.72				
	NT2RP1000416	9.92	9.55	19.32	9.98	6.1	6.11	5.35	8.54	6.35				
	NT2RP1000418	5.47	5.48	8.66	6.04	3.54	3.99	4.07	4.75	2.23				
	NT2RP1000420	22.79	21.95	33.37	16.39	16.84	13.03	9.29	9.88	11.08	*		-	
10	NT2RP1000434	13.91	15.33	17.37	9.5	9.94	9.87	16.13	15.05	16.89	**		-	
	NT2RP1000439	25.29	35.1	29.04	26.55	26.88	19.87	29.22	23.74	23.38				
	NT2RP1000443	20.2	15.54	27.59	12.04	8.66	7.66	5.79	6.64	10.23	*	*	-	-
	NT2RP1000447	8.08	15.72	18.03	9.49	4.04	7.9	4.98	7.69	18.07				
	NT2RP1000448	3.96	4.08	14.01	5.34	1.34	3.27	1.83	3.09	2.48				
	NT2RP1000451	4.39	7.13	7.39	6.28	2.08	2.05	2.16	2.95	1.76	*		-	
15	NT2RP1000458	7.7	7.09	11.38	7.06	4.06	2.49	4.41	2.72	2.94	*		-	
	NT2RP1000465	3.88	6.63	8.77	6.04	2.34	3	3.32	3.69	3.78				
	NT2RP1000468	10.59	11.91	14.42	9.41	9.31	7.51	8.52	6	7.58	*	*	-	-
	NT2RP1000470	5.79	2.68	10.41	5.27	1.59	4.02	1.42	2.51	2.5				
	NT2RP1000477	92.32	107.32	97.76	75.25	82.97	57.94	30.6	83.57	85.18	*		-	
	NT2RP1000478	15.37	19.72	31.58	11.53	8.67	8.82	5.74	7.51	6.16		*	-	
20	NT2RP1000481	26.31	30.78	29.03	24.24	26.58	11.39	8.04	10.78	12.11	**		-	
	NT2RP1000493	20.43	14.94	23.34	16.46	8.49	8.21	13.43	15.08	18.68				
	NT2RP1000513	46.32	35.58	52.07	60.82	43.2	48.5	36.73	37.16	31.64				
	NT2RP1000522	31.35	42.08	42.76	21.13	28.37	20.27	18.09	8.09	12.01	*	**	-	-
	NT2RP1000533	3.26	4.01	2.16	4.33	1.36	1.15	1.34	1	0.91	*		-	
	NT2RP1000544	15.77	9.6	19.95	8.16	5.29	4.62	4.57	9.59	8.27	*		-	
25	NT2RP1000547	6.82	3.97	11.11	7.34	2.91	3.12	1.8	4.52	2.98				
	NT2RP1000551	7.57	6.25	20.26	6.79	4.51	3.98	3.69	5.06	3.29				
	NT2RP1000567	7.5	7.79	12.24	11.47	5.44	5.98	2.72	3.24	2.13	*		-	
	NT2RP1000574	10.39	7.63	14.91	6.66	4.35	2.94	2.41	2.6	3.1	*		-	
	NT2RP1000577	6.72	6.64	7.37	6.99	2.31	2.08	2	3.14	1.89	**		-	
	NT2RP1000579	6.59	4.72	10.08	5.29	2.62	2.74	2.3	1.65	2.85	*		-	
30	NT2RP1000581	11.66	12.4	16.41	11.21	8.47	7.93	11.76	7.04	4.88				
	NT2RP1000593	11.3	18.39	27.51	6.2	5.55	5.03	1.75	7.5	4.02	*	*	-	-
	NT2RP1000604	24.51	26.11	37.01	27.05	13.43	25.97	13.69	26.68	23.37				
	NT2RP1000609	11.66	13.43	21.36	13.95	9.13	6.99	3.92	12.37	8.18				
	NT2RP1000613	17.95	20.62	29.59	15.28	14.93	12.95	12.7	12.66	9.93	*		-	
	NT2RP1000622	5.38	3.34	9.13	5.39	2.07	1.13	2.51	2.21	0.46				
35	NT2RP1000627	4.89	3.23	5.07	4.57	2.03	1.44	1.45	1.18	1.66	**		-	
	NT2RP1000629	8.08	7.69	11.36	7.12	3.71	2.97	5.95	2.89	3.59	*		-	
	NT2RP1000630	7.73	9.3	8.35	7.9	4.49	4.92	4.54	3.4	2.81	**		-	
	NT2RP1000639	15.2	13.58	16.62	7.62	5.84	8.02	4.83	7.17	5.87	**	**	-	-
	NT2RP1000640	7.36	9.59	17.46	9.53	6.74	8.49	5.28	6.04	7.68				
	NT2RP1000646	4.24	5.51	12.04	6.49	2.96	3.77	4.79	5.79	2.35				
40	NT2RP1000659	23.32	18.8	20.24	8.66	11.69	4.88	4.8	5.33	5.99	**	**	-	-
	NT2RP1000674	5.93	7.17	11.83	6.23	3.45	3.48	1.76	2.58	3.57	*		-	
	NT2RP1000677	14.6	24.32	25.71	10.51	12.87	7.3	5.18	10.1	7.28	*	*	-	-
	NT2RP1000679	49.27	69.62	36.95	29.7	31.01	14.48	34.6	17.18	10.04				
	NT2RP1000688	8.01	6.65	7.8	7.2	2.8	2.42	6.02	2.46	2.38	*		-	
	NT2RP1000689	32.73	30.66	30.93	19.81	8.55	17.9	17.63	24.88	21.39	*	**	-	-
45	NT2RP1000695	17.35	14.85	15.87	10.45	3.47	5.36	2.82	6.77	7.19	*	**	-	-
	NT2RP1000701	7.19	4.67	10.84	7.55	5.43	2.64	2.6	8.76	2.45				
	NT2RP1000702	7.2	10.25	10.38	5.65	2.25	1.64	3.18	3.24	2.79	*	**	-	-
	NT2RP1000713	7.21	6.46	7.91	6.66	2.57	3.12	4.23	3.89	1.7	*		-	
	NT2RP1000721	15.01	8.63	33.56	6.6	3.3	4.77	2.95	3.62	2.32				
	NT2RP1000730	34.22	37.43	42.87	27.42	15.22	17.38	23.72	15.85	27.4	*	*	-	-
50	NT2RP1000733	9.95	10.41	8.91	8.92	3.88	5.5	7.94	2.92	5.71	*		-	
	NT2RP1000738	1.62	1.49	4.67	5.56	1.13	3.01	1.45	2.92	1.28				
	NT2RP1000739	19.87	17.5	27.32	13.76	6.59	15.22	15.07	28.31	27.46				
	NT2RP1000740	10.94	13.28	13.49	8.2	7.61	6.64	6.43	11.92	8.57	**		-	
	NT2RP1000746	3.82	5.92	7.73	5.64	4.22	2.64	2.76	3.96	2.29				
	NT2RP1000750	12.79	9.22	13.66	7.67	4.28	4.14	7.19	4.29	4.16	*	*	-	-
55	NT2RP1000751	10.42	10.46	12.4	8.57	5.65	4.98	4.2	5.02	5.07	*	**	-	-
	NT2RP1000767	10.46	7.93	10.97	6.93	2.33	2.54	1.45	2.21	3.81	*	**	-	-
	NT2RP1000769	4.03	4.96	3.53	5.29	3.09	2.58	3.04	5.45	2.56				

Table 418

	NT2RM4002634	8.22	4.99	13.16	9.11	4.37	3.31	1.21	3.45	2.39			
	NT2RM4002636	6.59	6.68	10.88	5.63	3.08	3.09	2.42	3.32	2.29	*		-
5	NT2RP1000002	38.84	33.13	45.9	29.04	32.42	26.61	13.28	16.85	17.96	**		-
	NT2RP1000005	16.16	12.37	15.85	10.42	5.99	5.72	9.46	7.81	8.28	*	**	-
	NT2RP1000015	7.32	9.13	12.92	6.24	3.84	1.22	4.29	2.29	4.19	*		-
	NT2RP1000018	7.99	5.66	4.74	10.16	3.74	4.75	7.93	7.02	6.47			-
	NT2RP1000034	93.68	89.94	88.11	64.65	53.66	86	72.54	94.51	102.97			-
10	NT2RP1000035	18.17	13.62	22.35	15.17	8.64	11.35	7.31	9.51	10.39	*		-
	NT2RP1000040	16.4	12.21	23.34	12.38	8.64	8.09	5.04	9.39	4.11	*		-
	NT2RP1000042	3.77	4.47	8.66	6.24	2.47	2.4	1.5	1.51	0.43	*		-
	NT2RP1000048	11.21	13.33	14.61	8.93	8.14	5.51	8.58	7.29	5.06	*	*	-
	NT2RP1000050	5.73	12.28	7.67	5.34	2.29	5.33	2.26	2.09	1.81	*		-
	NT2RP1000056	13.77	17.86	17.27	9.41	8.14	4.91	6.92	3.33	2.85	**	**	-
15	NT2RP1000058	2.7	2.41	1.36	4.8	0.95	2.39	2.76	1.39	2.38			-
	NT2RP1000063	17.16	17.65	11.27	8.7	3.73	4.52	5.36	8.5	8.3	*	*	-
	NT2RP1000068	29.14	23.7	36.49	16.13	16.83	12.51	21.73	27.58	32.86	*		-
	NT2RP1000072	39.99	32.45	48.63	26.8	23.08	25.9	35.59	35.1	31.66	*		-
	NT2RP1000073	30.81	35.89	43.46	17.05	8.38	10.22	9.08	9.67	9.54	**	**	-
	NT2RP1000078	9.27	9.25	12.57	6.24	5.48	5.5	2.69	7.89	4.73	*	*	-
20	NT2RP1000079	5.69	6.1	5.36	5.63	2.36	2.02	2.99	2.23	1.19	**		-
	NT2RP1000080	8.49	8.44	8.35	6.53	2.81	2.65	4.1	3.05	2.07	*	**	-
	NT2RP1000086	8.5	8.4	9.89	12.4	14.28	6.92	7.93	6.49	7.97	*		-
	NT2RP1000087	90.2	68.37	62.22	84.59	56.96	88.52	51.31	53.27	70.97			-
	NT2RP1000089	42.88	33.03	33.53	24.16	25.57	29.27	22.12	38.13	29.4	*		-
	NT2RP1000090	36.01	34.5	36.9	42.34	38.64	45.67	26.7	26.11	29.29	*	**	+
25	NT2RP1000100	642.61	1542.2	1396.7	306.69	827.1	384.76	1509.9	896.88	763.21	*		-
	NT2RP1000101	35.43	98.15	94.63	31.12	33.53	50.26	29.26	23.54	17.24	*	**	-
	NT2RP1000111	7.76	7.95	10.33	5.26	2.18	1.26	2.31	1.62	2.57	*	**	-
	NT2RP1000112	23.37	50.82	42.64	24.98	11.61	20.15	18.5	13.38	13.37	*		-
	NT2RP1000124	25.62	22.22	19.79	7.84	2.34	7.4	4.52	4.07	4.19	**	**	-
	NT2RP1000125	10.72	8.67	12.07	7.14	2.95	7.79	4.48	7.44	4.25	*		-
30	NT2RP1000129	53.88	53.28	57.25	42.69	30.82	50.19	47.48	64.98	53.17			-
	NT2RP1000130	19.47	14.75	28.76	11.14	15.19	12.69	8.3	10.06	14.98			-
	NT2RP1000154	3.93	3.52	5.78	5.04	1.58	2.69	2	2.85	2.74			-
	NT2RP1000163	9.61	17.58	16	11.58	5.58	5.03	6.39	4.65	5.07	*		-
	NT2RP1000170	4.53	4.62	7.7	6.31	2.3	2.01	2.67	3.54	2.2			-
	NT2RP1000174	9.99	10.63	9.3	6.41	2.75	3.81	5.44	4.09	3.96	**	**	-
35	NT2RP1000181	14.26	14.61	16.73	10.84	15.44	12.79	10.24	5.66	5.53	**		-
	NT2RP1000191	12.96	10.86	7.92	16.4	11.07	15.31	6.1	13.72	9.91			-
	NT2RP1000202	24.16	14.94	24.74	18.17	11.59	9.79	7.42	9.13	15.3			-
	NT2RP1000239	9.53	5.6	7.62	9.29	2.89	11.84	8.48	12.01	11.59			-
	NT2RP1000243	6.22	4.43	4.68	7.85	5.29	6.41	2.34	3.31	5.03			-
40	NT2RP1000255	12.37	12.15	15.41	16.34	7.48	13.06	9.35	10.82	12.98			-
	NT2RP1000259	10.81	8.35	10.69	9.6	2.44	9.12	13.53	11.26	13.39			-
	NT2RP1000261	8.39	2.66	5.85	15.75	5.57	6.12	7.38	4.73	4.58			-
	NT2RP1000269	2.25	3.12	1.21	5.48	2.06	2.18	4.74	1.71	2.26			-
	NT2RP1000271	20.23	11.57	26.77	18.9	8.83	20.16	9.94	14.85	10.84			-
	NT2RP1000272	2.12	2.3	9.68	5.22	1.59	2.18	2.89	2.48	2.25			-
	NT2RP1000279	10.54	8.91	14.66	10.51	4.61	8.63	10.77	11.05	13.98			-
45	NT2RP1000290	6.06	4.1	11.1	13.63	4.43	7.63	6.81	6.19	8.17			-
	NT2RP1000293	3.83	4.19	7.55	5.54	1.62	3.86	2.95	1.13	3.16			-
	NT2RP1000331	4.37	2.83	13.34	5.34	2.19	3.45	2.73	6.07	2.66			-
	NT2RP1000333	10.16	8.8	16.01	10.64	5.56	6.94	8.71	5.87	8.68			-
	NT2RP1000336	18.67	16.33	24.52	12.77	10.82	14.23	17.03	16.82	19.7			-
	NT2RP1000347	3.46	4.07	7.92	6.45	3	4.58	4.14	3.03	3.73			-
50	NT2RP1000348	34.17	39.38	47.03	19.99	17.54	15.37	14.54	7.37	12.73	**	**	-
	NT2RP1000349	3.22	4.33	6.7	6.49	4.16	3.15	4.9	4.23	3.34			-
	NT2RP1000353	180.66	121.66	192.83	151.27	98.19	147.88	118.94	165.86	159.59			-
	NT2RP1000356	8.01	8.29	12.26	8.92	4.2	7.42	6.21	5.29	12.39			-
	NT2RP1000357	38.01	31.62	38.61	32.16	21.31	27.88	24.36	33.21	30.56			-
	NT2RP1000358	9.65	6.55	14.28	10.15	5.03	7.68	4.14	4.11	2.37	*		-
55	NT2RP1000360	11.08	7.25	16.85	18.78	9.48	8.41	8.09	9.21	9.08			-
	NT2RP1000363	37.26	30.39	52.28	41.73	39.62	39.87	33.79	24.41	33.64			-
	NT2RP1000376	6.26	6.27	9.53	5.76	4.97	3.76	4.87	4.58	3.63			-

Table 417

	NT2RM4002240	41.31	33.08	34.96	23.86	21.39	33.98	42.08	38.96	43.61			
	NT2RM4002251	3.39	3.69	6.47	4.89	3.61	2.4	2.93	2.15	17.6			
5	NT2RM4002256	4.28	3.3	5.05	4.37	2.61	2.38	2.35	2.33	1.46	*	-	-
	NT2RM4002262	25.4	21.79	26.58	14.04	9.23	12.13	20.41	14.34	24.05	**	-	-
	NT2RM4002266	32	29.71	28.68	24.62	15.02	21.58	34.67	32.96	37.69	*	-	+
	NT2RM4002276	14.45	13.71	15.67	14.53	13.36	16.53	16.37	10.08	9			
	NT2RM4002278	18.11	14.14	14.01	7.25	7.14	10.52	3.65	6.03	7.88	*	**	-
10	NT2RM4002281	9.29	5.59	12.79	12.53	4.74	10.06	6.18	9.79	8.22			
	NT2RM4002287	17.77	15.59	14.41	11.94	5.05	12.57	14.23	12.4	14.92			
	NT2RM4002294	5.99	6.14	6.14	5.51	2.38	5.99	3.64	2.35	2.91	**	-	-
	NT2RM4002298	64.75	62.55	58.47	42.58	28.72	43.54	69.57	64.59	61.46	**	-	-
	NT2RM4002301	3.57	3.23	3.63	6.49	2.3	3.07	3.59	2.99	3.21			
	NT2RM4002306	25.86	19.13	24.79	14.02	9.84	17.37	30.57	32.67	35.82	*	-	+
15	NT2RM4002323	7.61	6.27	5.98	7.59	3.26	7.4	8.04	6.9	4.8			
	NT2RM4002334	14.45	6.6	16.75	21.2	9.98	15.23	10.19	12.3	13.22			
	NT2RM4002339	36.85	25.91	51.52	34.79	27.9	40.82	21.13	20.38	29.16			
	NT2RM4002344	17.19	8.04	22.21	10.94	8.27	8.93	15.04	14.72	15.42			
	NT2RM4002345	7.58	5.39	12.54	5.63	3.79	3.52	2.31	4.45	2.41			
	NT2RM4002352	44.06	33.96	54.03	33.98	35.28	37.69	25.21	30.31	27.97			
20	NT2RM4002362	39.95	26.35	45.23	28.03	24.7	27.77	28.92	25.64	24.85			
	NT2RM4002373	9.44	6.99	7.1	8.69	3.38	4.42	5.88	3.57	3.37	*	-	-
	NT2RM4002374	0.94	1.53	2.98	5.8	0.19	1.91	2.19	2.71	2.57			
	NT2RM4002376	14.29	7.3	14.66	12.48	5.38	11.68	12.33	14.05	16.09			
	NT2RM4002383	5.57	4.67	12.54	5.61	3.39	2.94	3.24	5.28	5.45			
	NT2RM4002390	8.2	7.31	15.14	6.97	3.2	6.3	4.57	6.71	7.54			
25	NT2RM4002398	16.51	8.98	16.5	15.05	12.49	8.92	7.11	8.76	5.38			
	NT2RM4002409	3.91	1.86	5.55	4.87	1.55	2.44	2.22	2.93	0.87			
	NT2RM4002414	18.86	15.16	32.88	21.65	23.59	21.32	20.36	19.06	17.22			
	NT2RM4002439	3.96	4.99	8.08	7.18	3.61	4.94	4.85	4.1	2.07			
	NT2RM4002440	32.39	17.01	28.92	38.22	20.77	30.03	30.94	26.11	25.22			
	NT2RM4002446	10.16	7.87	12.29	14.29	6.56	8.04	5.21	8.63	33.19			
30	NT2RM4002450	5.69	5.77	10.55	7.51	2.92	4.53	1.81	3.53	7.42			
	NT2RM4002452	15.3	12.15	20.1	12.34	7.95	8.52	9.04	11.33	9.95			
	NT2RM4002457	5.93	4.98	9.2	5.51	3.34	2.74	3.42	3.45	3.16			
	NT2RM4002458	5.3	7.18	9	7.94	5.54	4.54	4.93	4.05	3.12			
	NT2RM4002460	5.82	9.3	10.61	8.18	6.55	6.42	6.02	6.26	3.91			
	NT2RM4002464	5.03	6.81	7.71	5.37	2.82	2.84	2.52	3.29	2.23	*	-	-
35	NT2RM4002479	40.28	37.05	46.08	41.61	24.81	36.24	37.84	34.54	34.16			
	NT2RM4002482	13.66	18.31	28.36	15.72	10.35	15.81	9.15	8.99	19.35			
	NT2RM4002489	15.44	12.03	20.84	12.49	8.35	8.73	7.84	10.26	14.04			
	NT2RM4002493	8.53	4.66	11.35	5.77	2.46	3.69	1.55	3.88	3.31			
	NT2RM4002499	16.96	15.44	26.91	11.24	8.53	18.09	18.09	16.97	14.06			
	NT2RM4002504	24.74	20.67	17.4	17.01	12.22	12.67	5.73	8.11	7.63	**	-	-
40	NT2RM4002506	9.92	11.26	12.04	6.45	4.68	4.22	2.88	4.99	2.98	**	-	-
	NT2RM4002510	17.14	20.34	27.62	13.7	8.64	11	8.3	5.68	5.29	*	-	-
	NT2RM4002527	5.27	5.32	5.24	9.73	2.73	4.8	7.63	6.1	4.65			
	NT2RM4002532	2.04	3.38	12.81	3.92	2.01	0.75	2.59	1.83	2.16			
	NT2RM4002534	6.24	4.2	12.1	6.74	2.8	3.26	1.86	3.47	2.45			
	NT2RM4002536	9.02	8.87	16.41	7.91	3.99	2.92	0.97	4.02	2.94	*	-	-
45	NT2RM4002554	6.47	7.38	8.92	4.87	2.87	2.29	2.04	1.83	1.15	*	**	-
	NT2RM4002558	10.72	10.21	14.55	8.7	6.67	5.1	5.89	7.82	7.9	*	*	-
	NT2RM4002565	7.12	9.02	9.26	6.37	4.46	3.35	5.55	4.05	5.46	*	*	-
	NT2RM4002567	8.35	6.05	12.48	6.62	4.6	3.2	2.87	5.5	6.18			
	NT2RM4002571	3.38	3.03	4.65	5.3	2.26	1.53	2.86	1.83	1.84			
	NT2RM4002572	3.42	1.92	7.29	5.02	1.66	0.9	0.78	0.74	1.09			
50	NT2RM4002577	17.78	12.61	20.74	8.52	7.02	10.13	3.18	8.47	5.97	*	*	-
	NT2RM4002583	26.66	37.71	48	24.28	17.87	13.18	6.06	13.66	11.34	*	*	-
	NT2RM4002584	24.55	35.28	41	25.45	15.38	14.04	7.11	11.26	5.72	**	-	-
	NT2RM4002593	13.18	11.32	14.62	7.42	4.85	4.9	3.83	4.05	2.43	**	-	-
	NT2RM4002594	17.08	15.66	20.52	11.9	10.96	5.12	3.63	4.19	2.45	*	**	-
	NT2RM4002604	8.24	6.88	8.88	5.84	3.74	2.95	2.85	0.99	1.52	*	**	-
55	NT2RM4002614	14.57	8.99	9.81	12.06	5.4	4.41	5.49	5.87	6.71	*	*	-
	NT2RM4002616	31.41	23.47	38.09	25.19	21.94	0.64	13.84	17.54	27			
	NT2RM4002629	6.22	6.78	13.92	7.63	3.69	4.81	3.87	5.8	5.63			

Table 416

	NT2RM4001885	71.31	49.62	81.98	53.98	43.04	45.93	68.83	64.57	48.87				
	NT2RM4001889	10.85	7.93	15.01	8.59	4.29	4.67	4.77	6.73	3.99				
5	NT2RM4001894	30.62	32.08	44.07	33.7	34.28	35.89	33.56	30.42	25.18				
	NT2RM4001897	8.76	10.37	11.73	7.79	4.55	4.65	3.35	3.47	2.33	*	**	-	-
	NT2RM4001899	4.19	5.18	10.36	7.76	5.5	4.54	4.99	5.17	3.45				
	NT2RM4001905	8.04	9.67	9.34	12.13	4.62	8.49	17.26	7.42	9.7				
	NT2RM4001922	10.52	7.5	13.46	10.13	4.31	4.77	4.26	7.33	9.75				
10	NT2RM4001930	42.04	27.93	35.21	26.34	18.3	24.57	35.05	37.8	24.49				
	NT2RM4001938	11.86	11.93	17.44	9.19	6.81	6.54	7.09	10.88	7.99	*		-	-
	NT2RM4001940	40.15	39.79	42.12	29.48	25.03	26.5	19.62	21.25	18.41	**	**	-	-
	NT2RM4001942	7.8	8.46	12.88	6.27	3.74	4.54	3.46	2.56	2.64	*		-	-
	NT2RM4001953	2.56	5.71	6.3	5.41	2.95	2.84	2.21	3.3	2.4				
	NT2RM4001965	9	9.88	11.74	9.82	12.01	7.33	7.4	10.04	8.64				
15	NT2RM4001966	51.68	52.74	61.28	70.45	40.89	59.05	88.01	53.06	72.22				
	NT2RM4001969	9.53	10.35	15.87	9.41	5.57	8.09	5.17	7.89	12.14				
	NT2RM4001974	12.36	21.98	16.69	16.74	8.35	12.5	5.35	17.32	22.51				
	NT2RM4001979	23.89	19.24	22.43	14.04	16.93	18.71	17.76	22.69	22.51				
	NT2RM4001980	94.16	97.18	94	64.27	69.09	81.77	75.44	72.18	71.63	*	**	-	-
	NT2RM4001984	9.67	8.39	11.58	7.27	4.78	4.84	2.35	4.35	2.8	*	**	-	-
20	NT2RM4001987	7.05	10.61	11.42	5.92	3.18	5.3	2.56	4.02	2.33	*	**	-	-
	NT2RM4002013	15.54	16.72	17.68	12.71	8.41	7.52	18.32	10.81	11.68	*		-	-
	NT2RM4002018	18.51	18.01	18.38	23.63	14.06	15.67	18.72	13.95	13.04				
	NT2RM4002033	6.2	4.97	12.78	7.92	4.07	6.05	2.33	3.53	3.27				
	NT2RM4002034	9.06	7.36	15.25	10.05	4.49	7.82	4.17	10.89	9.93				
	NT2RM4002044	29.35	20.58	32.58	18.82	16.22	17.19	21.2	22.06	24.1				
25	NT2RM4002047	20.66	17	25.07	17.46	15.74	9.49	16.1	15.14	16.08				
	NT2RM4002054	31.19	30.54	36.08	20.52	17.55	11.13	16.47	15.97	19.8	**	**	-	-
	NT2RM4002055	21.78	18.7	21.85	12.08	11.37	8.76	14.04	16.74	17.09	**	*	-	-
	NT2RM4002059	10.36	12.77	15.52	6.8	7.53	3.99	4.33	4.04	2.53	*	**	-	-
	NT2RM4002061	9	11	9.24	10.93	6.82	5.33	6.65	4.88	4.34	**		-	-
	NT2RM4002062	97.19	68.62	99.9	109.04	77.98	163.08	52.62	69.12	71.01				
30	NT2RM4002063	8.61	5.55	11.75	8.31	3.06	3.95	4.25	4.81	4.94				
	NT2RM4002066	7.96	8.56	22.27	12.72	22.89	4.97	2.38	10.48	2.89				
	NT2RM4002067	10.87	18.12	21.45	11.17	10.75	7.5	4.23	11.4	5.44				
	NT2RM4002073	34.67	29.49	32.35	16.42	15.53	13.85	5.51	8.53	5.15	**	**	-	-
	NT2RM4002074	13.13	16	13.48	9.5	9.68	8.27	2.81	4.16	5.74	**	**	-	-
	NT2RM4002075	12.66	9.69	13.41	10.12	6.41	6.32	4.7	4.84	4.77	**		-	-
35	NT2RM4002076	5.05	4.42	4.84	7.24	1.99	2.02	1.56	2.18	1.6	**		-	-
	NT2RM4002078	61.79	41.41	62.13	44.43	23.08	49.45	46.13	69.55	79.59				
	NT2RM4002081	12.47	11.63	21.52	11.3	5.46	8.36	4.38	5.75	7.06	*		-	-
	NT2RM4002082	28.78	23.31	28.71	31.25	27.4	16.48	17.58	15.02	15.49	**		-	-
	NT2RM4002093	8.61	9.51	12.26	11.79	5.58	6.25	5.84	7.44	5.28	*		-	-
	NT2RM4002109	5.91	6.93	10.41	8.11	5.84	4.18	3.3	3.14	2.63	*		-	-
40	NT2RM4002115	9.41	7.3	7.03	6.27	4.66	1.89	3.74	1.98	2.7	**		-	-
	NT2RM4002118	37.45	58.41	50.21	20.19	26.02	17.58	13.66	5.39	8.55	*	**	-	-
	NT2RM4002128	8.86	5.7	4.74	7.73	3.31	3.74	6.06	4.03	5.02				
	NT2RM4002137	10.72	7.46	13.29	9.5	3.57	7.97	5.86	8.61	6.51				
	NT2RM4002139	4.03	2.83	9.32	5.97	2.53	3.54	2.11	4.86	1.91				
	NT2RM4002140	13.52	11.7	18.78	14.09	10.76	8.24	10.97	13.55	10.92				
45	NT2RM4002145	9.85	17.52	15.66	7.49	7.31	6.12	9.29	8.25	6.8	*		-	-
	NT2RM4002146	53.03	50.09	60.28	48.49	40.37	39.91	51.33	53.96	59.3	*		-	-
	NT2RM4002161	42.85	50.59	52.56	30.41	24.15	28.58	27.02	24.86	21.9	**	**	-	-
	NT2RM4002174	10.01	10.09	12.11	7.53	4.92	3.2	3.7	2.29	3.16	*	**	-	-
	NT2RM4002178	12.55	13.35	13.21	15.19	8.36	11.5	9.67	9.09	6.99	**		-	-
	NT2RM4002180	29.73	19.51	29	18.7	7.96	18.84	13.92	22.12	26.06				
50	NT2RM4002185	43.79	41.64	55.35	22.67	17.47	26.66	26.11	42.39	45.09	**		-	-
	NT2RM4002189	7.67	6.89	10.29	5.96	5.23	2.74	5.92	2.8	1.53	*		-	-
	NT2RM4002194	41.77	35.89	39.22	19.41	14.19	21.77	29.88	28.52	30.53	**	*	-	-
	NT2RM4002198	65.16	103.48	83.23	24.68	28.75	36.65	24.56	34.17	38.68	**	*	-	-
	NT2RM4002205	10.02	12.63	11.47	6.5	3.78	3.22	4.61	2.19	6.22	**	**	-	-
	NT2RM4002213	21.88	23.22	24.09	13.45	7.91	12.62	20.89	17.99	21.66	**		-	-
55	NT2RM4002216	7.77	7.03	5.98	10.01	6.36	6.33	6.06	4.45	5.36				
	NT2RM4002226	41.57	37.46	37.46	67.73	42.88	64.51	42.92	35.21	34.67				
	NT2RM4002237	7.28	7.73	10.06	4.52	3.54	3.6	2.94	7.15	4.84	**		-	-

Table 415

	NT2RM4001565	12.7	17.39	26.89	14.14	12.4	12.69	7.45	6.92	4.69	*	-
	NT2RM4001566	17.21	11.12	20.03	22.75	14.87	13.43	10.31	12.88	15.08		
5	NT2RM4001569	3.45	2.79	9.43	4.65	1.24	1.1	2.19	2.38	1.74		
	NT2RM4001579	7.75	7.8	9.32	7.49	3.8	1.35	2.94	2.54	2.36	**	-
	NT2RM4001582	6.95	5.09	5.63	5.76	1.73	2.34	1.66	1.19	2.1	**	-
	NT2RM4001589	113.7	167.05	103.64	41.43	57.6	46.23	27.72	15.73	11.4	*	**
	NT2RM4001592	6.17	5.42	14.24	6.52	2.18	4.41	4.13	6.42	4.93		
10	NT2RM4001594	386.49	313.56	285.69	287	185.32	238.47	160.25	299.69	275.08		
	NT2RM4001597	6.23	6.27	13.54	6.73	3.64	4.25	2.96	4.23	3.88		
	NT2RM4001605	16.38	18.33	22.25	14.56	11.19	8.87	10.5	12.13	9.23	*	*
	NT2RM4001609	23.54	15.9	22.4	13.71	9.55	9.27	6.4	7.54	6.99	*	**
	NT2RM4001610	10.55	8.52	11.74	9.35	5.13	4.46	6.11	6	5.63	*	-
	NT2RM4001611	14.22	14.7	15.89	10.73	7.27	6.47	6.73	5.19	3.38	**	**
15	NT2RM4001618	13.74	16.13	14.23	8.24	4.82	5.02	4.84	2.49	3.4	**	**
	NT2RM4001622	5.3	5.16	10.27	7.37	3.16	3.66	2.49	3.88	1.85		
	NT2RM4001624	7.85	5.37	11.44	5.27	1.81	3.96	3.07	2.98	3.67	*	-
	NT2RM4001625	8.28	12.41	10.38	15.36	14.48	2.63	7.15	9.31	10.39		
	NT2RM4001629	14.67	21.52	22.9	19.11	13.87	22.31	12.28	12.44	12.55	*	-
	NT2RM4001632	325.71	457.93	338.97	242.41	290.06	370.51	594.61	333.67	482.68		
20	NT2RM4001642	5.97	4.92	5.47	4.64	2.19	2.19	0.94	2.25	0.91	*	**
	NT2RM4001647	18.42	12.21	17.82	18.58	14.27	17.66	13.87	14.03	11.49		
	NT2RM4001650	27.91	31.97	47.33	36.24	30.14	33.6	22.2	16.78	26.83		
	NT2RM4001662	12.58	10.55	10.99	18.39	7.75	15.61	11.98	12.75	8.09		
	NT2RM4001666	7.92	11.39	9.77	4.29	5.85	7.46	4.49	4.14	4.36	*	**
	NT2RM4001670	25.82	19.71	20.86	17.87	11.22	17.38	19.89	23.01	24.94		
25	NT2RM4001682	8.89	10.21	10.73	7.71	7.31	3.19	3.94	4.22	3.18	**	-
	NT2RM4001710	8	9.89	13.67	6.48	2.86	3.01	1.39	4.38	1.64	*	*
	NT2RM4001712	6.13	6.94	9.77	4.06	2.96	2.15	2.17	4.16	1.89	*	*
	NT2RM4001714	18.44	20.46	18.31	8.4	3.92	5.06	6.84	3.44	4.9	**	**
	NT2RM4001715	22.24	32.3	32.67	17.97	6.58	16.35	15.07	7.8	8.58	*	*
	NT2RM4001727	17.83	12.1	16.99	14.9	8.57	19.16	13.54	13.69	14.31		
30	NT2RM4001731	9.18	5.56	7.2	4.83	1.9	3.44	2.33	4	3.02	*	*
	NT2RM4001735	15.03	8.65	13.3	10.72	9.69	6.1	6.95	9.42	9.18		
	NT2RM4001739	3.31	4.05	7.1	6.51	1.7	2.71	2.2	2.1	1.76		
	NT2RM4001741	5.03	6.39	6.03	4.45	2.85	1.9	3.44	3.9	2.72	*	**
	NT2RM4001746	27.18	35.45	35.48	20.34	15.62	12.62	17.33	16.68	21.57	**	*
	NT2RM4001754	47.9	62.7	56.46	79.4	39.53	40.99	48.7	37.13	39.56		
35	NT2RM4001757	4.77	10.74	3.2	6.77	1.73	2.53	2.83	4.79	2.31		
	NT2RM4001758	8.89	8.75	6.1	8.32	3.51	5.38	5.88	8.51	9.88		
	NT2RM4001768	3.37	0.36	1.12	5.36	1.88	2.21	0.35	1.5	0.9		
	NT2RM4001775	6.53	3.93	5.59	8.91	3.87	5.25	4	3.54	3.16		
	NT2RM4001776	28.42	28.92	28.25	33.08	15.74	17.3	19.26	20.43	19.52	**	-
	NT2RM4001783	4.96	4.55	4.74	5.37	2.6	3.32	2.42	2.8	2.84	**	-
40	NT2RM4001793	14.95	13.94	19.52	14.23	7.85	12.31	14.46	13.31	17.08		
	NT2RM4001810	23.52	24.77	30.47	26.07	19.33	20.51	34.17	38.53	24.96		
	NT2RM4001813	9.51	6.59	9.34	11.22	3.7	8.5	12.3	12.03	14.87	*	+
	NT2RM4001818	4.65	1.67	8.39	7.01	2.88	5.12	5.55	5.88	5.41		
	NT2RM4001819	80.16	35.79	68.43	71.44	40.61	80.22	41.62	48.42	47.06		
45	NT2RM4001823	10.42	6.03	12.1	10.8	5.46	5.65	5.68	8.09	6.63		
	NT2RM4001828	3.57	2.82	6.95	5.66	1.65	1.21	2.42	1.92	2.31		
	NT2RM4001835	4.94	4.42	12.73	8.71	1.8	3.96	4.63	4.11	5.11		
	NT2RM4001836	5.12	3.81	8.12	5.95	1.63	1.85	2.61	3.45	2.12		
	NT2RM4001841	6.73	6.82	10.79	10.33	4.39	6.34	6.7	6.45	7.77		
	NT2RM4001842	2.89	4.82	4.7	7.5	2.83	2.4	3.82	3.29	2.69		
	NT2RM4001843	48.57	28.12	59.83	46.07	40.73	45.97	28.16	28.11	29.49		
50	NT2RM4001856	14.1	14.11	19.35	11.87	5.04	7.86	7.47	6.5	8.37	*	*
	NT2RM4001858	20.02	12.7	24.11	14.43	7.61	10.18	17.09	15.78	20.13		
	NT2RM4001861	14.08	10.48	18.84	13.72	9.53	5.84	12.28	18.05	13.36		
	NT2RM4001863	27.15	15.67	30.03	15.24	18.38	17.87	16.63	16.94	11.79		
	NT2RM4001865	8.03	9.87	15.29	7.44	8.43	5.34	6.68	5.7	6.95		
	NT2RM4001869	12.34	16.35	21.36	13.19	9.35	9.55	18.86	14.54	12.88		
55	NT2RM4001873	6.51	9.79	10.95	7.15	2.28	3.17	4.21	4.29	2.39	*	-
	NT2RM4001876	13.28	10.02	39.33	14.58	13.74	11.65	8.08	13.34	13.31		
	NT2RM4001880	22.41	18.46	28.54	18.91	14.09	15.45	13.74	18.99	19.02		

Table 414

	NT2RM4001100	10.34	6.97	14.94	8.81	5.31	8.81	7.75	11.53	7.75			
	NT2RM4001116	6.46	5.43	9.66	8.08	5.44	5.9	5.27	4.35	5.99			
5	NT2RM4001119	4.41	2.95	6.53	4.88	4.54	2.33	3.03	2.56	2.24			
	NT2RM4001140	6.58	8.81	10.22	5.74	2.39	2.78	3.27	2.64	2.47	*	**	-
	NT2RM4001148	77.43	82.01	87.89	109.01	92.69	80.91	71.63	54.14	60.54	*		-
	NT2RM4001151	25.3	23.87	22.4	14.14	12.5	10.51	24	19.61	17.03	**		-
	NT2RM4001155	7.7	10.96	9.45	14.26	6.55	9.63	8.68	7.88	7.8			
	NT2RM4001157	4.69	2.46	3.77	5.35	1.74	4.67	1.85	2.85	2.34			
10	NT2RM4001160	10.13	5.87	8.44	9.04	3.06	4.73	3.63	6.09	4.88			
	NT2RM4001163	8.69	6.99	8.41	6.85	4.09	5.89	3.57	3.78	3.99	**		-
	NT2RM4001187	15.55	10.66	18.41	9.39	4.2	6.99	4.74	5.75	3.81	*	*	-
	NT2RM4001191	6.6	5.61	4.99	6.79	10.14	6.02	8.39	4.7	8.34			
	NT2RM4001200	2.58	2.46	3.46	4.38	3.72	1.51	1.91	2.35	0.33			
15	NT2RM4001203	6.43	5.54	5.53	5.95	1.89	1.97	3.96	4.4	1.73	*		-
	NT2RM4001204	9.94	10.43	11.48	7.05	1.89	2.44	3.49	2.67	2.11	*	**	-
	NT2RM4001217	24.69	17.41	30.8	23.78	12.47	18.57	27.41	26.41	29.34			
	NT2RM4001245	156.95	84.48	149.16	130.37	116.53	157.67	151.03	141.35	178.46			
	NT2RM4001247	4.65	4.81	11.8	4.49	3.5	3.11	6.79	3.17	2.29			
	NT2RM4001256	9.61	7.67	12.78	9.59	5.91	6.14	7.73	8.06	8.51			
20	NT2RM4001258	70.47	64.71	70.52	44.98	46.66	50.08	53.38	45.51	40.98	**	**	-
	NT2RM4001267	5.89	4.4	10.4	6.42	2.33	2.85	4.9	3.94	4.52			
	NT2RM4001273	1.34	2.22	3.89	5.02	1.41	1.99	4	2.5	2.16			
	NT2RM4001281	8.06	5.61	14.03	10.91	4.33	7.34	4.05	7.62	5.35			
	NT2RM4001286	51.19	32.28	36.36	29.23	18.59	18.64	13.09	20.76	31.46			
	NT2RM4001290	8.38	5.81	13.36	7.76	4.34	4.65	5.33	4.36	7.29			
25	NT2RM4001309	98.2	68.64	139.85	90.19	105.33	108.01	85.7	73.37	76.26			
	NT2RM4001313	6.16	7.7	10.44	8.55	3.72	5.52	2.76	5.94	3.03			
	NT2RM4001316	10.67	13.16	19.79	13.68	8.94	6.99	6.58	5.71	5.93	*		-
	NT2RM4001320	20.75	22.22	27.83	18.9	15.02	17.67	25.16	19.45	25.01			
	NT2RM4001321	3.2	9.27	5.51	7.64	4.65	3.21	6.77	5.74	7.05			
	NT2RM4001325	6.99	10.29	13.71	11.04	4.07	7.04	5.14	7.06	21.4			
30	NT2RM4001333	11.2	7.57	11.93	8.81	7.22	5.3	2.36	6.09	14.33			
	NT2RM4001340	59.12	52.84	78.91	69.02	66.75	67.91	59.71	47.75	38.99			
	NT2RM4001344	6.4	4.63	10.13	7.69	4.29	5.37	3.9	5.77	3.72			
	NT2RM4001347	414.8	327.64	302.6	368.33	392.69	323.06	333.75	373.98	353.9			
	NT2RM4001357	6.26	7.97	8.82	6.78	6.02	4.63	3.22	2.89	2.9	**		-
	NT2RM4001360	8.63	8.04	10.78	8.59	6.1	3.5	4.75	7.61	3.39			
35	NT2RM4001371	15.56	11.85	10	11.21	16.5	13.92	36.9	13.45	6.6			
	NT2RM4001377	21.16	11.62	19.22	16.27	7.45	12.65	8.88	10.18	9.45			
	NT2RM4001382	16.22	9.22	22.97	11.49	7.37	7.19	3.28	8.65	16.08			
	NT2RM4001384	4.53	5.9	14.13	5.13	2.13	2.71	2.62	2.27	2.98			
	NT2RM4001400	10.43	9.23	12.46	8.14	5.92	4.37	4.37	6.39	3.72	*	**	-
	NT2RM4001409	4.67	5.55	8.19	6.09	3.26	2.83	3.17	3.27	3.39			
40	NT2RM4001410	6.89	14.26	12.51	8.64	5.38	3.51	3.98	6.53	3.24			
	NT2RM4001411	25.9	30.99	34.06	31.33	32.6	28.51	31.72	28.52	32.83			
	NT2RM4001412	825.39	1226.8	1011.2	1004.1	804.98	983.25	1210.2	810.05	943.27			
	NT2RM4001414	12.79	6.6	16.28	6.35	2.72	5.16	2.14	4.73	3.09	*		-
	NT2RM4001436	63.2	35.96	69.06	56.83	54.08	67.38	42.65	35.42	49.83			
	NT2RM4001437	26.18	27.89	39.17	20.31	15.52	18.05	16.35	28.68	22.74	*		-
45	NT2RM4001444	3.41	3.61	9.9	6.98	3.02	1.89	2.57	2.59	2.26			
	NT2RM4001454	19.23	20.25	21.89	16.73	15.4	12.09	15.71	18.45	13.57	*	*	-
	NT2RM4001455	5.88	19	6.35	4.26	2.07	3.65	1.94	1.57	2.98			
	NT2RM4001483	8.69	16.03	11.64	7.61	4.41	5.78	3.96	3.81	3.42	*		-
	NT2RM4001489	12.59	12.07	8.74	5.56	6.34	2.86	4.94	5.52	3.23	*	**	-
	NT2RM4001495	36.43	24.24	20.4	12.41	16.18	17.63	5.62	16.36	11.58			
50	NT2RM4001499	22.29	18.15	30.48	16.01	14.05	14.71	4.23	11.08	11.85	*		-
	NT2RM4001515	6.81	5.91	14.18	6.91	2.64	4.03	2.9	4.57	2.36			
	NT2RM4001519	8.21	4.24	9.85	5.65	1.45	0.78	1.08	3.15	1.34	*		-
	NT2RM4001522	8.19	6.53	12.02	6.15	2.52	1.81	2.28	3.64	1.62	*		-
	NT2RM4001523	73.84	75.98	71.57	35.61	37.16	20.7	26.91	23.43	33.3	**	**	-
	NT2RM4001550	10.75	6.21	9.98	6.43	6.16	2.2	5.6	2.68	8.03			
55	NT2RM4001553	11.94	9.8	11.96	9.12	5.14	4.64	3.81	2.93	3.19	*	**	-
	NT2RM4001554	151.86	72	141.69	164.83	103.78	171.98	80.79	76.13	147.24			
	NT2RM4001557	51.14	42.32	58.99	56.35	49.65	76.92	34.69	30.81	38.74	*		-

Table 413

	NT2RM4000733	9.25	6.15	14.75	8.98	6.63	6.77	4.25	7.21	11			
	NT2RM4000734	6.83	6.23	12.29	7.58	4.56	3.67	3.75	5.75	9.29			
5	NT2RM4000741	13.34	10.33	15.73	9.84	9.36	7.54	6.59	7.87	3.9	*		-
	NT2RM4000744	885.53	451.3	1241	1383.2	1506.5	707.31	777.95	717.55	894.48			
	NT2RM4000749	51.3	49.81	57.77	49.46	44.79	42.49	59.41	56.25	54.64			
	NT2RM4000751	10.01	10.59	16.02	7.97	7.75	5.05	5.19	5.27	4.04	*		-
	NT2RM4000752	14.18	16.62	13.54	10.44	6.55	5.76	8.04	11.07	5.33	*	*	-
10	NT2RM4000760	56.56	27.61	44.38	43.74	22.69	40.29	50.15	65.89	60.41			
	NT2RM4000761	19.9	12.95	27.34	11.59	8.66	7.67	5.75	12.58	17.06			
	NT2RM4000764	28.14	13.4	30.32	14.61	9.8	20.16	18.5	24.69	24.19			
	NT2RM4000768	91.29	73.54	83.67	77.4	70.39	93.31	63.42	64.16	68.6	*		-
	NT2RM4000778	8.24	6.22	8.88	7.38	4.94	5.13	2.54	2.81	2.48	**		-
	NT2RM4000779	13.14	10.85	20.24	8.82	8.68	7.51	2.45	7.69	5.22	*		-
15	NT2RM4000787	15.66	16.22	17.04	13.21	10.45	11.66	11.77	12.92	11.34	**	**	-
	NT2RM4000790	2620.9	2248.3	2489	2090	3251.6	2752.2	3676.2	1775.7	1256.3			
	NT2RM4000795	7.92	4.23	13.52	7.6	3.13	5.92	6.03	8.23	6.4			
	NT2RM4000796	26.78	15.24	27.73	18.41	12.57	18.03	14.45	19.92	38.07			
	NT2RM4000798	7.01	8.1	18.9	8.2	4	6.33	3.86	7.47	8.07			
	NT2RM4000800	5.35	6.72	10.01	6.15	1.63	3.53	2.61	2.89	3.11	*		-
20	NT2RM4000813	11.25	10	13.48	6.95	4.25	5.74	3.76	4.39	3.13	**	**	-
	NT2RM4000820	26.13	26.55	28.07	22.17	20.99	17.91	22.8	23.35	20.32	**	*	-
	NT2RM4000827	10.03	14.9	13.91	7.34	7.52	5.34	7.29	6.84	3.6	*	*	-
	NT2RM4000830	12.07	12.08	13.71	8.9	8.35	9.39	11.27	7.81	7.47	**	*	-
	NT2RM4000833	11.24	7.17	16.3	8.57	3.24	6.92	2.5	5.31	5.24			
	NT2RM4000841	12.98	15.96	14.24	9.21	12.52	7.68	4.07	5.69	5.39	**		-
25	NT2RM4000846	9.5	7.99	19.32	10.93	5.53	8.29	6.17	10.68	8.48			
	NT2RM4000848	7.99	8	13.35	6.98	3.66	2.79	2.7	4.77	2.57	*		-
	NT2RM4000852	14.87	14.63	15.93	7.89	7	3.89	3.95	4.67	2.59	**	**	-
	NT2RM4000855	6.4	5.44	7.14	5.82	2.45	2.19	1.77	2.66	1.11	**		-
	NT2RM4000859	16.62	15.67	16.78	13.33	8.27	6.7	12.27	10.96	12.51	*	**	-
	NT2RM4000868	19.91	23.16	23.62	14.39	12.92	10.23	4.82	6.2	3.98	**	**	-
30	NT2RM4000870	13	13.65	19.75	13.98	14.72	8.66	5.75	8.62	12.86			
	NT2RM4000878	41.44	32.18	43.25	14.78	13.02	16.56	3.72	8.51	8.81	**	**	-
	NT2RM4000882	15	11.7	22.42	13.18	7.01	7.9	5.18	5.53	4.82	*		-
	NT2RM4000887	41.97	33.25	51.62	36.36	23.39	28.87	18.02	26.87	25.16	*		-
	NT2RM4000895	26.8	20.11	33.05	17.45	16.12	7.46	11.88	6.63	10.98	*		-
	NT2RM4000897	9.87	8.53	11.07	8.99	4.54	3.34	4.8	3.39	4.04	**		-
35	NT2RM4000901	10.96	12.11	11.53	7.56	7.1	4.42	5.41	4.23	5.77	**	**	-
	NT2RM4000950	9.78	8.98	10.35	8.13	5.43	3.19	2.51	2.31	1.79	*	**	-
	NT2RM4000965	28.19	32.79	35.21	15.2	12.26	17.85	6.17	10.47	12.16	**	**	-
	NT2RM4000971	97.81	77.57	101.74	132.86	95.27	155.13	87.73	92.85	108.16			
	NT2RM4000979	6.96	6.06	14.84	8.87	6.06	5.77	4.88	5.81	4.09			
	NT2RM4000987	998.76	831.62	1103.5	625.74	1156.8	778.33	1010.1	1262.3	668.26			
40	NT2RM4000988	20.87	13	27.71	19.07	13.82	14.2	14.36	15.32	14.84			
	NT2RM4000991	41.92	23.13	35.75	27.46	31.48	28.64	21.97	22.75	24.51			
	NT2RM4000992	10.32	6.68	8.53	6.07	2.23	2.13	3.02	2.16	2.78	*	**	-
	NT2RM4000996	22.09	29.46	18.01	12.6	11.82	10.41	9.81	3.95	3.92	*	*	-
	NT2RM4000997	28.21	30.93	25.95	19.16	13.92	19.47	16.5	21.62	24.11	**	*	-
	NT2RM4001001	7.63	8.6	15.15	9.35	5.11	5.88	3.96	7.75	5.13			
45	NT2RM4001002	459.98	441	503.26	494.01	440.37	522.33	521.26	628.75	590.54	*		+
	NT2RM4001016	19.09	20.3	24.95	12.05	10.46	13.91	10.42	12.83	14.8	*	*	-
	NT2RM4001025	6.64	7.24	14.2	7.08	5.08	3.26	5.04	3.42	2.59			
	NT2RM4001027	4.69	5.86	4.64	4.07	1	1.53	1.18	1	1.23	*	**	-
	NT2RM4001032	13.63	13.78	6.47	6.13	6.41	2.73	4.98	2.77	2.21	*		-
	NT2RM4001047	17.2	23.07	17.27	11.83	13.43	8.74	14.05	4.88	2.78	*	*	-
50	NT2RM4001049	14.8	12.01	16.43	12.47	5.42	11.65	9.35	13.4	14.12			
	NT2RM4001051	23.07	23.96	19.46	18.43	11.09	9.95	3.77	14.4	12.67	*	*	-
	NT2RM4001052	9.72	6.17	11.79	8.32	6.45	2.86	8.57	9.7	6.4			
	NT2RM4001053	8.35	8.48	10.92	7.64	6.3	5.31	6.88	7.16	3.68			
	NT2RM4001054	15.68	17.03	15.78	11.07	4.7	7.63	9.63	9.71	11.5	*	**	-
	NT2RM4001059	5.24	11.23	8.09	5.5	2.85	2.53	2.48	2.39	3	*		-
55	NT2RM4001071	23.88	28.79	33.58	18.66	9.28	10.59	17.49	15.69	16.83	*	*	-
	NT2RM4001084	12.62	12.59	13.06	12.4	5.97	9.54	14.16	9.85	15.23			
	NT2RM4001092	7.67	7.08	7.41	8.26	5.02	4.62	2.05	6.72	4.2			

Table 412

	NT2RM4000306	39.39	31.22	45.45	18.05	17.89	26.02	12.1	14.22	20.93	*	**	-	-
	NT2RM4000307	10.91	9.3	17.74	9.83	6.51	5.04	4.55	5.59	9.36				
5	NT2RM4000309	6.25	6.98	13.33	6.03	3.16	3.04	0.89	2.68	1.25	*		-	
	NT2RM4000313	8.25	9.27	15.88	8.4	4.79	4.48	3.48	4.24	4.11	*		-	
	NT2RM4000318	7.74	12.3	19.97	11.61	7.91	4.9	8.74	11.58	10.87				
	NT2RM4000324	7.02	4.79	7.97	5.66	2.78	1.11	2.3	1.61	0.69	**		-	
	NT2RM4000326	9.61	6.92	8.36	6.7	2.41	2.62	3.8	2.41	1.32	**		-	
	NT2RM4000327	10.62	7.04	10.88	10.24	4.86	9.97	11.64	7.37	9.72				
10	NT2RM4000344	28.12	43.85	37.76	23.98	18.35	22.38	19.35	25.83	27.75	*		-	
	NT2RM4000349	54.86	44.05	74.07	45.06	38.46	49	53.73	62.48	76.26				
	NT2RM4000354	3.83	4.95	10.6	6.31	1.75	1.8	2.77	2.77	2.79				
	NT2RM4000356	5.03	3.65	9.35	5.42	1.79	2.58	1.32	3.38	0.81				
	NT2RM4000366	170.18	124.74	186.28	108.07	122.86	98.86	143.05	122.68	104.4				
	NT2RM4000368	16.94	13.21	17.9	9.95	5.92	5.45	9.44	7.79	10.81	*	*	-	-
15	NT2RM4000373	32.64	29.04	37.96	31.16	21.65	24.87	33.11	24.62	25.87				
	NT2RM4000386	6.62	6.28	8.3	8.65	4.71	4.93	5.02	2.91	5.23				
	NT2RM4000395	10.95	13.82	15.83	10.78	7.59	9.02	5.57	7.71	10.22	*		-	
	NT2RM4000414	4.69	4.37	11.16	5.76	0.91	3.43	2.88	1.6	3.45				
	NT2RM4000417	11.27	10.89	8.84	6.8	4.15	6.41	8.68	3.94	4.92	*		-	
	NT2RM4000421	10.28	10.99	14.3	8.17	4.31	2.7	2.33	2.02	4.78	*	**	-	-
20	NT2RM4000425	91.24	92.57	106.69	64.91	58.82	58.32	76.16	70.13	74.53	**	*	-	-
	NT2RM4000433	6.13	6.12	8.53	7.28	5.88	4.18	2.6	4.18	3.51	*		-	
	NT2RM4000436	18.9	19.36	27.32	20	14.51	11.77	15.16	14.79	15.69				
	NT2RM4000444	18.5	21.83	23.5	27.88	22.92	21.75	22.28	13.95	20.52				
	NT2RM4000457	13.16	19.12	21.52	10.27	6.05	7.18	7.2	10.64	8.19	*	*	-	-
	NT2RM4000471	11.07	11.03	12.38	8.14	9.49	4.57	3.84	6.85	3.58	**		-	
25	NT2RM4000472	19.66	22.31	17.4	12.44	15.19	12.8	11.64	22.04	27.4	*		-	
	NT2RM4000486	10.68	18.04	19.73	11.41	9.93	10.39	10.79	12.22	9.89				
	NT2RM4000490	14.19	17.43	15	9.61	4.76	5.97	7.87	5.19	5.16	**	**	-	-
	NT2RM4000496	4.23	6.24	5.23	3.18	0.15	0.92	1.52	0.44	1.12	*	**	-	-
	NT2RM4000505	107.79	139.53	149.56	114.44	75.32	107.4	136.82	107.52	141.57				
	NT2RM4000511	73.01	84.21	82.3	77.22	39.83	88.6	83	54.31	48.26				
30	NT2RM4000514	15.58	17.06	20.66	15.1	9.63	16.07	11.41	15.96	14.62				
	NT2RM4000515	15.97	18.43	19.43	7.56	4.12	3.43	3.77	6.97	4.37	**	**	-	-
	NT2RM4000517	351.24	469.1	440.94	138.38	381.29	140.48	205.12	297.92	317.1	*		-	
	NT2RM4000520	3.56	5.44	6.75	6.2	2.43	4.29	3.06	2.99	1.91				
	NT2RM4000531	7.21	6.94	9.49	7.75	2.88	4.52	5.61	3.99	4.45	*		-	
35	NT2RM4000532	6.55	5.58	4.11	7.27	2.1	1.72	3.22	3.22	3.25	*		-	
	NT2RM4000533	5.69	4.73	4.92	6.83	1.35	2.87	3.57	2.18	3.55	*		-	
	NT2RM4000534	4.83	3.29	2.48	5.97	1.61	1.76	3.52	3.71	3.07				
	NT2RM4000563	23.61	15.27	21.44	28.24	15.01	45.63	27.71	31.71	29.82	*		+	
	NT2RM4000566	9.4	6.65	10.24	13	4.81	13.26	8.88	9.31	8.58				
	NT2RM4000568	24.52	13.96	20.25	24.28	11.25	22.83	7.78	15.36	11.01				
40	NT2RM4000585	3.91	1.91	3.09	7.76	1.73	4.77	2.79	4.1	3.39				
	NT2RM4000587	7.59	6.74	8.76	10.89	3.52	5.69	7.38	7.94	10.18				
	NT2RM4000590	2.8	2.23	4.43	6.82	1.35	3.31	2.85	3.67	2.59				
	NT2RM4000593	8.21	5	10.19	8.56	3.29	5.21	6.43	4.7	4.53				
	NT2RM4000595	4.79	4.42	3.36	7.11	6.02	3.43	4.16	2.82	3.87				
	NT2RM4000603	5.4	2.57	9.66	6.3	2.52	2.39	2.28	2.67	2.18				
45	NT2RM4000611	28.57	18.91	36.37	25.67	15.39	24.58	19.91	19.95	26.79				
	NT2RM4000616	9.69	7.13	13.2	7.2	3.11	4.07	2.46	3.02	2.66	*		-	
	NT2RM4000621	3.69	1.04	7.35	6.04	1.52	1.89	3.47	3.26	2.31				
	NT2RM4000648	7.93	5.22	9.62	8.39	3.12	4.35	6.48	5.01	6.06				
	NT2RM4000649	4.41	3.44	7.74	6.28	2.95	1.85	3.85	1.71	2.41				
	NT2RM4000658	5.46	6.14	9.39	8.84	2.67	4.42	4.63	4.69	4.24				
50	NT2RM4000661	15.52	9.28	15.4	13.03	7.61	7.72	5.33	7.59	7.97	*		-	
	NT2RM4000673	13.87	8.93	21.18	9.61	6.09	5.58	4.94	6.04	10.95				
	NT2RM4000674	33.39	14.79	28.45	15.54	18.31	16.46	9.71	12.48	12.12				
	NT2RM4000689	13.32	8.68	20.09	9.6	6.2	5.32	9.33	7.16	5.74				
	NT2RM4000698	3.25	4.45	8.82	7.51	4.65	2.16	4.55	4.72	3.77				
	NT2RM4000700	9.2	12.82	18.94	9.98	10.71	10.66	5.56	6.11	5.56	*		-	
55	NT2RM4000701	37.49	38.63	57.82	37.22	35.87	38.99	50.37	52.92	48.89				
	NT2RM4000712	26.37	34.47	28.93	18.91	14.79	13.59	9.02	8.85	10.91	**	**	-	-
	NT2RM4000717	24.65	10.07	20.07	21.94	7.33	18.21	17.18	23.49	18.21				

Table 411

	NT2RM2002764	5.49	12.15	7.39	6.52	3.1	7.78	2.89	4.03	5.76				
	NT2RM2002772	29.76	33.25	38.82	20.56	12.45	9.21	4.96	16.05	9.09	**	**	-	-
5	NT2RM2002811	16.39	25	25.21	9.15	15.71	3.55	6.58	8.61	10.62	*	*	-	-
	NT2RM2002818	20.34	18.56	21.22	13.83	9.54	15.73	17.66	20.72	25.61	*	*	-	-
	NT2RM2002879	25.35	30.35	32.28	9.31	5.54	8.46	5.48	6.4	6.89	**	**	-	-
	NT2RM2002979	45.6	67	65.04	29.24	39.73	11.12	45.71	18.88	50.67	*	*	-	-
	NT2RM2002981	6.39	5.25	5.04	5.13	2.8	1.9	2.21	1.27	2.8		**	-	-
	NT2RM2002995	4.58	2.61	3.97	7.89	1.25	3.74	3.35	2.92	4.03			-	-
10	NT2RM2003031	4.98	5.14	6.7	6.97	3.27	6.19	2.14	4.35	4.33			-	-
	NT2RM2003042	32.29	28.58	51.11	28.69	7.79	11.66	9.6	10.82	7.31	*	*	-	-
	NT2RM2003044	5.51	2.76	2.92	6.07	1.88	3.64	2.44	2.91	3.92			-	-
	NT2RM2003090	15.51	15.57	25.03	16.18	8.25	19.49	18.74	20.12	15.75			-	-
	NT2RM2003095	23.07	26.27	29.36	22.31	14.66	13.94	29.67	24.13	25.38	*	*	-	-
	NT2RM2003116	19.38	32.85	27.63	12.52	9.52	10.31	6.49	8.73	7.15	*	**	-	-
15	NT2RM2003222	4.03	3.37	4.48	6.23	1.12	2.42	2.91	3.67	3.87			-	-
	NT2RM2003224	23.91	19.9	24.52	18.14	9.42	15.36	24.57	16.25	13.82	*	*	-	-
	NT2RM2003250	119.92	69.2	166.74	153.09	107.43	107.61	63.52	77.81	74.98			-	-
	NT2RM2003258	14.27	11.09	23.88	11.45	8.67	5.92	4.04	5.75	7.4			-	-
	NT2RM2003262	13.63	6.96	21.6	7.07	3.47	2.7	6.15	5.4	5.15			-	-
20	NT2RM4000023	16.25	15.39	24.54	14.62	6.88	9.36	9.34	7.69	6.58	*	*	-	-
	NT2RM4000024	5.57	7.1	11.59	9.02	4.21	3.87	3.82	4.08	2.65			-	-
	NT2RM4000027	4.73	3.54	9.85	6.85	4.18	2.53	4.65	2.99	2.04			-	-
	NT2RM4000030	5.72	8.1	11.8	8.56	5.42	3.94	5.02	4.04	3.02			-	-
	NT2RM4000033	5.95	4.77	7.64	13.08	3.88	7.38	10.23	6.96	6.85			-	-
	NT2RM4000034	14.03	8.95	17.36	14.71	7.52	11.06	7.66	12.11	11.07			-	-
	NT2RM4000046	5.52	5.71	10.27	5.43	2.75	2.16	4.42	4.99	3.5			-	-
25	NT2RM4000052	3.66	3.61	9.75	4.88	2.31	3.26	1.99	3.02	2.4			-	-
	NT2RM4000054	24.72	16.37	22.59	16.92	16.99	12.56	13.57	15.39	11.37	*	*	-	-
	NT2RM4000061	2.45	2.46	7.19	5.03	3.91	3.49	2.41	2.69	0.75			-	-
	NT2RM4000074	109.94	111.99	162.79	132.85	134.31	127.91	155.38	114.81	153.05			-	-
	NT2RM4000085	16.76	26.39	27.87	18.47	15.63	12.58	12.15	6.78	9.96	*	*	-	-
30	NT2RM4000086	5.66	6.66	7.88	8.93	4.39	5.02	8.75	6.18	6.29			-	-
	NT2RM4000100	33.67	23.83	35.82	31.94	18.51	25.58	27.68	38.43	48.27			-	-
	NT2RM4000101	24.08	19.83	27.89	18.1	9.03	12.06	10.92	15.22	16.58	*	*	-	-
	NT2RM4000102	31.73	28.52	43	29.96	36.82	28.28	11.54	34.43	29.82			-	-
	NT2RM4000104	12.56	10.41	15.15	10.13	9.81	7.37	8.35	10.19	9.76			-	-
	NT2RM4000115	5.71	9.33	11.32	11.36	6.79	5.37	7.68	7.13	8.99			-	-
35	NT2RM4000129	2.55	5.58	5.12	5.02	4.24	3.57	3.26	2.78	3.18			-	-
	NT2RM4000139	5.53	8.52	11.67	6.45	6.29	5.11	4.73	4.33	5.31			-	-
	NT2RM4000149	3.74	4.09	4.62	8.9	2.03	3.09	4.1	3.48	3.35			-	-
	NT2RM4000155	17.15	10.36	17.5	14.41	8.06	9.43	4.52	7.7	6.79	*	*	-	-
	NT2RM4000156	43.23	34.65	52.17	33.06	31.74	35.59	32.83	34.2	32.88			-	-
	NT2RM4000167	5.13	4.59	12.54	4.2	2.19	0.82	1.69	2.03	1.71			-	-
40	NT2RM4000169	116.9	136.33	190.06	211.72	166.35	130.17	146.08	152.04	180.53			-	-
	NT2RM4000191	10.04	6.72	12.84	9.1	7.11	6.9	7.89	7.38	7.91			-	-
	NT2RM4000187	3.31	6.64	7.83	5.98	3.1	2.95	2.4	2.51	3.17			-	-
	NT2RM4000198	13.19	12.13	17.78	14.45	8.75	9.44	12.48	10.4	9.17			-	-
	NT2RM4000199	3.72	4.46	6.47	8.34	4.51	5.35	6.27	5.09	6.45			-	-
	NT2RM4000200	3.27	5.32	11.6	5.66	2.52	4.69	3.52	3.66	3.02			-	-
45	NT2RM4000202	7.07	4.86	13.64	7.53	1.78	3.26	4.51	5.82	5.86			-	-
	NT2RM4000210	5.45	5.44	15.45	6.67	3.23	1.92	1.86	2.7	2.32			-	-
	NT2RM4000215	8.8	9.1	12.69	7.49	4.69	2.59	3.73	4.19	3.48	*	**	-	-
	NT2RM4000220	21.3	15.96	19.47	12.48	9.78	7.44	10.31	10.33	6.87	*	**	-	-
	NT2RM4000229	4.29	6.1	7.3	4.96	2.64	1.79	2.82	2.26	3.55	*	*	-	-
	NT2RM4000231	12.18	22.17	22.16	12.3	13.79	7.12	11.29	7.48	7.64	*	*	-	-
50	NT2RM4000233	21.08	19.41	23.05	29.48	16.53	15.89	23.49	16.8	20.56			-	-
	NT2RM4000244	7.04	6.33	12.22	6.72	4.83	3.85	2.65	5.84	4.45			-	-
	NT2RM4000251	8.01	7.64	14.53	7	8.84	3.91	3.55	5.44	5.4			-	-
	NT2RM4000255	8.99	6.74	14.46	7.34	5.4	4.93	3.56	3.66	2.79	*	*	-	-
	NT2RM4000265	13.77	11.08	20.66	10.54	8.41	4.23	9.98	13.42	10.55			-	-
	NT2RM4000283	213.58	220.77	174.14	136.23	205.5	145.83	158.43	132.77	224.43			-	-
55	NT2RM4000284	85.27	81.74	108.73	78.21	81.89	45.3	89.45	84.2	83.7			-	-
	NT2RM4000290	13.22	11	18	10.81	6.15	3.95	7.71	6.59	6.44	*	*	-	-
	NT2RM4000295	9.32	6.67	7.61	11.47	5.8	4.29	6.42	5.34	6.78			-	-

Table 410

	NT2RM2001941	6.04	4.24	11.07	8.57	3.76	4	3.69	5.77	3.42			
	NT2RM2001950	10.88	9.24	13.32	9.74	3.68	5.38	5.27	9.54	12.08			
5	NT2RM2001952	5.22	5.96	18.37	6.97	3.67	2.12	3.56	5.93	8.5			
	NT2RM2001976	43.9	27.85	44.8	42.85	38.56	26.62	16.38	23.06	20.86	*		-
	NT2RM2001982	3.43	5.03	6.82	6.99	3.67	2.56	3.51	3.09	3.63			
	NT2RM2001983	12.18	15.12	21.04	15.68	19.01	6.03	9.47	6.45	4.73	*		-
	NT2RM2001984	20.37	33.54	33.12	16.47	10.56	10.16	8.25	7.86	3.8	*	**	-
	NT2RM2001989	3.86	5.39	6.78	8.98	3.51	2.63	4.78	3.3	2.04			
10	NT2RM2001996	16.29	18.4	29.01	16.49	15.54	12.47	7.55	13.49	15.15	*		-
	NT2RM2001997	11.71	10.41	18.16	7.7	4.63	5.51	3.14	5.18	8.39	*		-
	NT2RM2001998	9.82	8.18	17.59	8.92	5.29	6.42	2.82	8.96	6.36			
	NT2RM2001999	18.02	20.28	27.06	10.57	9.9	5.15	3.81	3.98	3.66	*	**	-
	NT2RM2002003	50.92	53.27	55.93	27	36.02	21.81	18.72	19.18	19.35	**	**	-
	NT2RM2002004	4.18	6.17	7.32	5.72	2.78	2.44	2.58	3.1	3.84			
15	NT2RM2002009	12.24	20.57	21.82	12.13	12.36	8.91	7.8	3.92	5.72	*		-
	NT2RM2002014	2.73	6.1	5.88	8.11	3.54	6	6.1	6.78	3.31			
	NT2RM2002019	125.13	124.6	184.05	89.22	90.25	54.05	33.48	56.11	68.43	*	*	-
	NT2RM2002029	13.94	31.19	27.2	28.59	19.23	17.84	11.48	35.39	33.05			
	NT2RM2002030	16.61	18.25	22.39	8.8	11.56	5.36	5.58	12.47	9.07	*	*	-
20	NT2RM2002034	58.72	73.27	50.5	59.63	56.44	27.9	21	20.77	37.34	*		-
	NT2RM2002049	21.26	22	23.61	14.68	15.64	11.74	6.05	9.95	7.32	**	**	-
	NT2RM2002055	3.27	6.4	6.36	5.47	2.96	2.6	3.63	3.29	2.55			
	NT2RM2002072	40.82	52.88	39.93	24.05	29.78	18.81	18.47	15.25	18	*	**	-
	NT2RM2002088	82.31	80.35	97	73.51	51.16	68.26	90.62	55.31	84.87			
	NT2RM2002091	22.22	21.11	34.44	9.76	9.27	5.82	2.71	4.2	5.17	*	**	-
25	NT2RM2002100	12.27	9.78	20.49	8.79	6.55	5.99	3.91	6.96	6.5			
	NT2RM2002109	4.21	5.64	16.28	7.68	3.73	3.18	3.74	2.46	4.42			
	NT2RM2002126	27.51	39.25	34.46	33.04	40.35	13.28	17.62	8.06	21.42	*		-
	NT2RM2002128	4.47	5.67	11.09	4.57	2.71	1.5	2.22	2.89	1.62			
	NT2RM2002129	22.51	19.86	17.11	13.35	16.48	7.76	5.96	8.88	6.92	**	**	-
	NT2RM2002142	23.72	27.99	22.69	11.44	14.47	7.07	5.7	5.27	5.78	**	**	-
30	NT2RM2002144	5.53	4.42	5.98	7	2.78	3.75	3.22	1.93	3.5	*	*	-
	NT2RM2002145	12.68	17.57	21.16	8.81	4.88	6.47	2.77	6.22	7.99	*	*	-
	NT2RM2002153	10.64	8.88	12.47	8.77	5.52	6.62	3.59	3.26	3.47	**	**	-
	NT2RM2002163	3.34	3.58	10.5	5.65	2.59	1.58	0.8	1.56	1.92			
	NT2RM2002170	9.23	12.24	23.46	10.21	7.83	3.14	3.4	4.81	4.17			
	NT2RM2002178	9.54	6.98	11.63	5.56	4.58	2.92	3.63	2.57	1.48	*	**	-
35	NT2RM2002179	8.2	10.06	9.02	7.04	6.34	2.72	2.79	2.76	2.85	**	**	-
	NT2RM2002270	5.65	6.66	8.96	6.13	3.31	1.8	2.35	2.99	1.98	*		-
	NT2RM2002326	8.26	5.94	6.26	7.55	3.63	3.3	2.53	3.14	5.2	*		-
	NT2RM2002337	9.91	15.68	18.49	12.85	7.23	4.54	5.02	6.94	5.77	*		-
	NT2RM2002339	6.62	10.73	17.92	7.67	6.02	5.59	3.93	4.03	4.49			
	NT2RM2002345	12.28	11.64	16.5	11.35	8.87	6.37	4.6	5.99	5.18	**		-
40	NT2RM2002368	11.9	12.43	21.01	8.78	6.6	5.66	4.97	7.35	7.34	*		-
	NT2RM2002381	5.04	4.54	7.61	5.17	3.68	1.33	2.23	2.48	3.79			
	NT2RM2002424	6.19	9.87	12.12	6.06	3.81	1.42	2.55	2.13	2.39	*		-
	NT2RM2002450	5.83	4.84	6.17	9.5	2.45	2.17	3.24	2.41	1.89	**		-
	NT2RM2002482	6.57	3.99	6.22	7.86	2.37	2.81	5.1	2.33	2.52			
	NT2RM2002492	115.88	159.06	153.05	67.65	86	93.3	108.8	133.63	153.04	*		-
45	NT2RM2002575	8.81	10.86	19.91	10.34	8.51	6.85	6.71	8.36	8.38			
	NT2RM2002580	12.81	11.63	17.51	7.68	4.03	7.15	4.84	5.41	5.92	*	**	-
	NT2RM2002592	30.1	60.89	57.77	16.85	24.42	15.38	23.91	11.59	15.71	*	*	-
	NT2RM2002608	107.02	108.16	125.35	69.13	83.62	75.26	88.4	51.95	98.01	**		-
	NT2RM2002615	21.13	24.19	16.26	10.72	11.75	7.36	11.09	7.92	5.92	*	*	-
	NT2RM2002622	37.43	55.62	40.03	34.45	40.44	21.34	16.57	11.49	8.97	**		-
50	NT2RM2002630	17.23	13.55	15.45	18.01	9.28	15.57	19.85	13.91	16.46			
	NT2RM2002634	10.84	11.13	17.05	9.62	2.19	5.41	7.89	7.53	7.31			
	NT2RM2002645	19.66	20.09	27.47	17.63	14.57	10.2	9.86	18.1	25.3			
	NT2RM2002646	40.87	51.15	69.9	15.35	22.28	15.25	17.48	14.11	15.62	*	*	-
	NT2RM2002647	50.8	59.26	54.64	17.81	19.82	20.32	24.11	18.36	23.98	**	**	-
	NT2RM2002652	15.86	23.36	17.16	10.87	6.59	8.39	7.1	6.49	11.22	*	*	-
55	NT2RM2002692	6.96	5.59	6.84	5.88	3.32	1.36	2.92	2.3	3.51	**	**	-
	NT2RM2002721	65.28	100.72	114.83	36.62	27.49	27.32	87.26	32.69	72.44	*		-
	NT2RM2002748	96.93	107.15	111.07	119.35	46.49	100.69	117.84	48.85	124.45			

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	NT2RM2001664	5.02	5.48	16.95	8.08	4.37	8.84	2.47	6.43	6.78				
	NT2RM2001668	15.06	19.03	25.63	5.79	6.13	6.44	2.42	3.93	2.92	*	**	-	-
5	NT2RM2001670	5.32	7.34	15.89	6.82	2.18	2.28	5.04	5.37	3.96				
	NT2RM2001671	8.87	8.86	12.78	9.4	4.31	5.72	3.03	4.34	3.98	**		-	
	NT2RM2001675	5.39	3.86	9.7	4.63	1.07	1.86	1.29	1.4	0.96	*		-	
	NT2RM2001681	10.15	3.72	7.87	6.68	2.43	1.75	3.17	2.29	1.77				
	NT2RM2001685	5.99	4.42	4.61	4.4	0.95	1.7	1.76	0.44	1.52	**		-	
10	NT2RM2001688	7.35	10.19	21.86	10.5	8.43	4.5	8.27	2.69	4.73				
	NT2RM2001695	21.84	15.61	27.76	16.05	12.1	18.2	12.98	24.27	27.96				
	NT2RM2001696	11.41	12.78	17.34	10.47	4.47	6.07	3.84	8.22	8.43	*		-	
	NT2RM2001698	4.46	5.04	9.55	6.62	3.5	2.86	2.2	3.3	2.08				
	NT2RM2001699	9.41	7.54	15.28	7.73	5.45	4.44	4.63	5.76	3.45				
	NT2RM2001700	6.01	3.92	15.92	7.19	8.76	1.93	1.65	2.78	1.4				
15	NT2RM2001704	15.56	11	13.6	7.65	5.94	5.33	7.22	3.63	5.28	**	**	-	-
	NT2RM2001706	17.58	13	15.93	12.4	7.58	8.26	9.21	8.02	9.01	*	**	-	-
	NT2RM2001714	6.57	6.52	7.48	5.63	2.25	2.43	2.36	2.21	2	*	**	-	-
	NT2RM2001716	8.11	6.76	8.16	6.9	3	2.28	1.75	4.87	3.64	*		-	
	NT2RM2001718	5.27	4.14	10.17	6.69	1.77	2.4	3.07	2.63	3.19				
	NT2RM2001723	7.52	14.9	19.44	7.99	8.79	1.56	2.66	6.02	4.35				
20	NT2RM2001727	13.96	15.14	20.23	10.17	9.39	6.64	16.21	10.41	9.88	*		-	
	NT2RM2001730	19.9	22.32	24.32	11.37	7.04	10.08	9.63	10.59	8.89	**	**	-	-
	NT2RM2001738	24.21	32	34.82	11.52	20.69	8.94	10.25	6.51	7.37	*	**	-	-
	NT2RM2001743	11.04	12.3	13.73	7.4	4.32	2.84	7.29	5	5.26	**	**	-	-
	NT2RM2001753	33.97	33.52	45.06	27.22	20.28	23.67	19.89	16.73	21.61	*	*	-	-
	NT2RM2001755	5.43	4.23	7.05	6	2.04	4.11	2.32	3.16	2.92	*		-	
25	NT2RM2001760	42.97	46.84	32.99	27.92	15.82	19.33	23.92	44.69	49.12	*		-	
	NT2RM2001765	3.06	3.62	6.89	4.79	2.41	2.93	4.69	3.78	2.64				
	NT2RM2001767	156.3	123.32	123.27	161.08	199.81	185.95	99.49	113.99	123.4	*		+	
	NT2RM2001768	10.21	7.85	24.02	6.47	5.49	3.96	2.75	3.87	3.14				
	NT2RM2001771	8.46	12.44	14.25	6.18	3.66	4.08	3.36	4	3.1	*	**	-	-
	NT2RM2001778	7.68	9.54	13.49	7.21	3	4.06	5.69	4.22	4.13	*		-	
30	NT2RM2001782	8.45	16.26	12.43	10.97	4.91	8.55	7.19	5.67	4.31				
	NT2RM2001784	2.33	3.38	2.94	5.28	0.42	1.26	1.88	3.54	2.58				
	NT2RM2001785	26.54	25.99	26.3	14.79	12.05	14.71	11.42	23.99	14.87	**		-	
	NT2RM2001792	9.85	6.48	8.81	6.44	5.86	3.8	3.68	3.95	3.88	*		-	
	NT2RM2001795	6.29	3.46	12.45	6.74	1.68	4.68	1.95	2.66	2.03				
	NT2RM2001797	5.8	5.87	12.89	5.74	3.15	4.35	4.29	3.75	2.07				
35	NT2RM2001800	8.98	9.98	12.02	6.5	3.87	3.21	3.89	2.41	4.66	*	**	-	-
	NT2RM2001803	9.13	11.59	12.8	14.78	5.7	4.76	5	4.99	6.43	**		-	
	NT2RM2001805	5.23	4.93	4.93	5.77	2.81	1.86	2.38	1.02	1.76	**		-	
	NT2RM2001806	9.59	8.37	10.47	8.03	2.61	4.16	1.45	2.5	1.61	**		-	
	NT2RM2001813	4.64	2.78	4.37	7.6	3.08	5.69	2.05	4.02	2.1				
	NT2RM2001814	4.12	2.59	2.69	6.32	1.65	3.13	2.04	1.23	1.1	*		-	
40	NT2RM2001818	2.04	2.24	0.99	4.82	0.99	0.95	0.43	0.73	1.2				
	NT2RM2001823	3.33	1.51	3.32	5	0.06	3.46	1.75	2.14	2.76				
	NT2RM2001825	18.28	14.86	18.88	21.81	10.23	18.33	13.27	20.99	16.7				
	NT2RM2001832	8.66	8.97	9.59	7.39	2.65	5.19	2.03	4.44	3.08	*	**	-	-
	NT2RM2001839	39.76	56.2	55.7	17.75	23.24	41.75	69.27	61.03	74.92				
	NT2RM2001840	16.73	9.83	12.67	16.44	10.4	10.16	11.51	19.3	18.18				
45	NT2RM2001851	13.19	13.13	16.5	15.12	5.82	6.1	7.33	10.63	8.57	*		-	
	NT2RM2001856	24.09	22.55	30.11	14.9	11.35	11.17	11.7	10.3	11.91	**	**	-	-
	NT2RM2001867	4.7	5.27	10.74	7.68	2.21	3.72	3.8	4.17	5.58				
	NT2RM2001869	150.4	263.47	450.85	333.27	292.69	360.02	366.03	265.55	360.35				
	NT2RM2001879	4.5	3.92	9.65	7	2.4	3.74	3.05	3.17	3.3				
	NT2RM2001883	0.5	1.23	3.35	4.6	1.17	2.06	2.26	1.81	2.31				
50	NT2RM2001886	5.2	8.33	7.95	8.22	4.29	6.46	4.42	3.66	6				
	NT2RM2001887	10.27	7.46	19.23	10.21	4.91	6.33	4.19	5.05	5.45				
	NT2RM2001896	1909.2	1304.9	1706.8	669.09	517.88	706.49	508.04	455.61	977.52	**	*	-	-
	NT2RM2001902	4.57	4.04	12.43	5.26	2.22	2.22	1.58	3.18	2.3				
	NT2RM2001903	63.33	63.93	80.39	57.98	54.4	31.36	56.94	50.98	47.6	*		-	
	NT2RM2001930	4.16	7.96	10.97	7.26	3.74	5.15	3.87	4.57	4.22				
55	NT2RM2001935	6.31	7.33	13.94	7.14	3.83	4.76	3.89	3.34	2.47				
	NT2RM2001936	4.31	7.8	13.64	6.57	5.23	5.41	6.76	3.21	3.62				
	NT2RM2001939	1.41	2.6	6.02	7.8	3.89	3.31	3.06	4.67	3.29				

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	NT2RM2001126	10.89	14.19	16.02	9.38	4.33	6.54	7.81	4.58	5.85	*	*	-	-
	NT2RM2001131	35.31	23.37	27.48	39.22	24.25	32.19	17.26	18.93	15.18	*	*	-	-
5	NT2RM2001141	23.07	16.72	24.48	15.15	8.42	15.3	16.57	25.01	24.73			-	-
	NT2RM2001152	7.28	6.36	6.81	7.22	2.67	4.03	3.72	4.85	2.92	**	**	-	-
	NT2RM2001177	16.43	13.33	22.76	9.09	9.55	10.29	11.03	12.82	14.78	*	*	-	-
	NT2RM2001194	18.95	35.2	33.45	15.63	10.75	13.96	9.18	10.24	10.12	*	*	-	-
	NT2RM2001195	22.22	32.77	41.71	19.89	14.02	10.87	14.76	17.07	21.12	*	*	-	-
10	NT2RM2001196	12.18	14.33	14.56	8.23	2.8	2.96	2.65	3.74	5.12	**	**	-	-
	NT2RM2001201	112.26	151.54	152.39	113.21	71.67	129.08	182.51	124.84	110.23			-	-
	NT2RM2001221	15.16	10.19	12.98	12.97	6.39	8.39	4.13	7.92	7.64	*	*	-	-
	NT2RM2001238	7.02	4.76	7.61	7.38	2.47	4.7	2.6	3.8	3.66	*	*	-	-
	NT2RM2001243	28.44	25.47	29.19	20	12.3	9.83	7.01	16.89	8.25	*	**	-	-
	NT2RM2001244	14.84	14.77	26.94	13.43	7.79	12.02	6.46	7.21	5.59	*	*	-	-
15	NT2RM2001247	149.17	156.12	167.83	262.88	153.51	292.21	159.18	102.66	137.39	**	**	-	-
	NT2RM2001256	12.96	15.03	17.12	8.24	5.02	5.55	4.85	4.32	6.34	*	*	-	-
	NT2RM2001269	10.95	13.1	12.53	11.07	7.62	4.32	9.04	9.38	5.97	*	*	-	-
	NT2RM2001278	10.38	10.85	11.45	12.61	4.01	9.04	14.51	13.94	16.53	**	+	-	-
	NT2RM2001291	8.24	5.62	12.69	7.3	4.57	5.55	4.22	4.08	3.96			-	-
	NT2RM2001294	64.54	65.77	111.27	63.88	47.1	58.52	73.02	74.33	85.75	*	*	-	-
20	NT2RM2001295	8.24	9.79	14.29	5.61	3.8	3.62	3.42	2.36	2.25	*	*	-	-
	NT2RM2001302	3.39	6.85	11.75	6.43	3.29	2.51	3.71	3.87	1.57			-	-
	NT2RM2001306	11.41	14.23	14.66	11.73	5.3	7.08	5.43	5.34	3.99	**	**	-	-
	NT2RM2001312	3.55	4.56	12.72	6.28	2.54	2.63	4.61	2.59	2.39			-	-
	NT2RM2001319	5.86	9.67	14.24	9.42	5.49	4.3	4.35	3.58	2.75			-	-
	NT2RM2001324	18.95	7.8	13.76	18.03	4.46	12.42	10.21	14.53	14.86			-	-
25	NT2RM2001345	28.65	21.37	24.71	26.89	8	18.65	5.12	14.99	21.21			-	-
	NT2RM2001360	17.92	12.3	24.16	13.81	8.68	10.67	6.5	8.95	13.99			-	-
	NT2RM2001370	10.29	7.05	19.66	11.56	5.81	4.36	5.18	4.87	5.67			-	-
	NT2RM2001391	7.29	4.99	11.62	6.8	2.82	5.45	4.23	6.86	5.98			-	-
	NT2RM2001393	12.33	6.71	15.2	11.88	7.78	6.57	5.84	9.03	6.99			-	-
	NT2RM2001420	8.12	9.09	13.52	7.85	4.31	5.05	3.66	4.14	3.54	*	*	-	-
30	NT2RM2001423	17.63	16.34	28	15.84	12.17	12	10.06	5.58	6.91	*	*	-	-
	NT2RM2001424	27.85	17.9	24.35	16.51	10.81	12.46	10.74	18.59	16.89	*	*	-	-
	NT2RM2001482	8.67	6.53	15.28	13.75	6.43	10.52	8.21	10.88	32.33			-	-
	NT2RM2001499	6.83	5.56	14.66	6.11	6.28	4.16	4.43	4.99	9.08			-	-
	NT2RM2001504	10.92	6.52	14.65	8.66	7.17	4.96	4.36	8.69	6.66			-	-
	NT2RM2001524	14.94	10.41	10.01	10.19	8.66	6.87	5.54	6.7	4.33	*	*	-	-
35	NT2RM2001530	3.04	4.86	7.01	5.84	1.95	3.09	2.61	3.78	1.3			-	-
	NT2RM2001533	31.06	33.52	48.05	22.65	24.5	20.86	46.4	35.86	44.09			-	-
	NT2RM2001540	13.31	12.86	17.79	14.39	9.31	10.73	9.68	7.65	7.91	*	*	-	-
	NT2RM2001544	6.59	3.7	12.06	6.73	1.95	5.1	3.34	3.2	4.05			-	-
	NT2RM2001547	17.98	17.35	32.25	17.06	12.78	13.61	10.07	14.99	28.06			-	-
40	NT2RM2001553	3.53	3.66	12.29	5.32	1.92	2.05	1.78	3.67	4.24			-	-
	NT2RM2001575	12.53	9.85	20.98	9.37	6.51	6.57	9.69	11.4	8.27			-	-
	NT2RM2001582	18.86	18.39	26.78	16.91	11.16	8.85	13.11	10.99	13.56	*	*	-	-
	NT2RM2001588	33.5	33.04	38.62	26.73	19.6	15.21	19.42	19.94	22.31	*	**	-	-
	NT2RM2001592	8.06	9.31	10.24	8.08	7.97	5.3	6.26	2.96	5.52	*	*	-	-
	NT2RM2001603	18.68	34.48	37.52	19.83	12.18	16.75	16.12	15.01	10.86			-	-
	NT2RM2001606	2.46	1.84	10.05	3.68	0.89	3.29	1.27	2.06	1.52			-	-
45	NT2RM2001611	8.26	11.08	21.04	8.21	4.42	4.61	5.24	5.95	7.18			-	-
	NT2RM2001613	31.23	41.23	53.28	26.42	31.28	14.27	16.42	46.76	35.82			-	-
	NT2RM2001626	7.76	5	11.59	5.79	3.19	1.08	2.56	3.93	1.62			-	-
	NT2RM2001632	12.51	10.55	19.26	10.94	8.41	4.21	3.48	2.87	3.67	*	*	-	-
	NT2RM2001633	9.38	12.28	13.2	7.51	9.17	4.14	3.74	4.79	3.89	**	**	-	-
	NT2RM2001635	8.99	10.05	10.56	5.7	4.03	2.66	6.33	4.93	5.79	**	**	-	-
50	NT2RM2001636	13.44	12.39	18.22	10.09	11.69	6.86	5.49	5.27	4.75	**	**	-	-
	NT2RM2001637	5.38	4.7	11.19	6.81	1.81	3.12	2.09	2.96	1.51			-	-
	NT2RM2001639	12.13	12.21	20.77	10.63	8.64	7	4.1	7.5	7.85			-	-
	NT2RM2001641	8.65	5.25	15	6.44	4.6	3.79	6.08	10.65	9.66			-	-
	NT2RM2001643	13.92	14.16	22.25	10.74	7	2.7	2.29	4.88	4.18	*	*	-	-
	NT2RM2001648	39.15	45.91	36.48	26.92	23.84	12.46	22.53	23.65	32.08	*	*	-	-
55	NT2RM2001652	12.26	10.35	14.74	6.98	5.99	3.53	3.25	2.26	3.06	*	**	-	-
	NT2RM2001659	16.4	10.95	13.51	8.64	9.21	1.42	3.18	2.99	3.04	**	**	-	-
	NT2RM2001660	10.08	9.77	12.95	6.57	5.46	2.86	2.55	2.98	2.6	*	**	-	-

Table 407

	NT2RM2000581	16.64	23.76	26.4	16.81	10.83	8.04	10.45	10.74	12.79	*	-
	NT2RM2000582	34.62	16.91	29.03	32.54	12.57	23.96	20.07	22	22.14		
5	NT2RM2000588	206.96	155.13	213.15	202.22	154.41	234.16	169.86	193.16	204.6	*	-
	NT2RM2000589	21.73	20.45	25.51	21.73	16.89	13.34	11.8	15.99	14.79	*	-
	NT2RM2000594	3.88	3.97	9.16	5.31	1.87	1.58	2.08	3.22	2.03		
	NT2RM2000599	23.3	37.43	42.86	41.55	34.51	20.59	19.28	20.01	29.72		
	NT2RM2000609	11.28	11.34	20.18	17.86	7.92	5.53	10.38	8.31	8.85		
	NT2RM2000612	12.76	13.34	15.09	12.28	9.14	7.94	10.91	11.84	11.22	*	-
10	NT2RM2000622	141.09	153.93	193.67	159.94	144.1	158.9	186.97	168.17	180.75		
	NT2RM2000623	10.1	4.92	12.51	17.89	6.76	12.96	6.19	8.79	10.66		
	NT2RM2000624	40.95	32.48	53.61	29.49	39.42	24.44	20.09	35.85	21.95		
	NT2RM2000632	4.22	16.93	13.59	10.55	4.72	5.77	3.71	9.75	42.27		
	NT2RM2000635	17.3	14.29	24.16	11.9	11.44	7.82	6.12	9.2	22.13		
	NT2RM2000636	8.64	6.1	10.92	8.1	4.79	4.03	4.23	7.12	5.02		
15	NT2RM2000639	12.38	11.39	15.28	10.66	6.59	7.7	5.72	6.03	7.49	* **	- -
	NT2RM2000649	21.7	22.5	30.17	18.52	14.23	12.19	12.61	12.94	13.29	* *	- -
	NT2RM2000658	32.82	43.08	46.86	26.73	41.37	32	30.61	19.44	22.2	*	-
	NT2RM2000660	14.51	13.27	29.92	14.04	12.55	16.72	10.13	15.37	13.11		
	NT2RM2000669	10.87	10.69	21.85	10.05	8.58	6.66	4.07	7.21	13.35		
20	NT2RM2000689	95.32	133.69	72.27	57.73	75.12	39.22	34.72	103.73	96.38		
	NT2RM2000691	8.29	7.99	12.66	7.17	5.02	1.98	1.67	3.91	3.3	*	-
	NT2RM2000714	22.27	18.36	27.51	17.39	20.26	19.54	14.42	10.68	9.4	*	-
	NT2RM2000718	7.47	11.22	8.73	7.49	4.21	3.74	6.42	4.3	4.39	*	-
	NT2RM2000732	30.31	39.24	31.84	16.98	20.8	19.07	19.55	19.14	20.81	** **	- -
	NT2RM2000735	9.9	11.53	13.44	9.07	5.38	4.22	4.93	3.86	5.02	* **	- -
25	NT2RM2000740	6.93	5.88	14.03	10.18	2.3	7.44	6.84	7.79	7.93		
	NT2RM2000743	6.68	6.8	14.3	6.91	4.28	3.65	5.06	3.56	3.44		
	NT2RM2000772	27.24	33.91	37.64	22.25	28.4	15.62	5.89	15.75	8.53	**	-
	NT2RM2000773	52.46	53.2	70.82	31.56	39.87	27.02	30.18	38.39	37.45	* *	- -
	NT2RM2000776	68.74	76.62	73.41	45.06	64.43	20.51	36.77	44.34	50.74	**	-
	NT2RM2000784	15.31	17.83	22.47	11.93	15.96	6.63	6.13	10.15	9.45	*	-
30	NT2RM2000795	27.14	19.23	31.28	19.3	19.19	9.44	22.75	21.68	23.6		
	NT2RM2000796	7.18	4.13	6.34	5.73	2.37	1.38	4.64	1.74	1.29		
	NT2RM2000798	220.92	157.62	241.6	287.45	173.39	313.04	164.61	215.06	234.9		
	NT2RM2000801	206.47	181.87	250.93	201.4	210	276.53	117.76	135.99	178.08		
	NT2RM2000821	19.71	21.75	22.88	13.91	8.23	8.65	3.33	5.35	4.94	** **	- -
	NT2RM2000829	32.72	37.9	42.47	37.61	27.12	23.67	12.82	16.2	13.46	**	-
35	NT2RM2000837	12.85	10.7	21.49	8.33	5.59	2.89	4.75	4.19	3.96	*	-
	NT2RM2000824	66.24	48.22	89.49	72.54	62.16	39.92	33.79	29.08	40.4		
	NT2RM2000930	48.55	63.69	52.84	46.58	51.6	35.79	27	25.97	35.7	**	-
	NT2RM2000937	31.8	58.4	47.42	22.54	35.77	15.38	12.6	7.67	4.38	**	-
	NT2RM2000939	6.62	8.62	14.67	7.32	3.05	3.47	2.42	2.17	3.83	*	-
	NT2RM2000942	734.65	670.2	631.32	344.25	467.07	301.87	150.27	478.62	381.55	** *	- -
40	NT2RM2000951	4.26	4.86	13.05	5.55	2.39	1.46	1.13	1.84	1.19		
	NT2RM2000952	9.67	43.75	19.11	12.99	13.49	9.68	6.64	9	6.45		
	NT2RM2000966	187.94	165.71	228.75	214.51	217.85	161.14	133.25	108.26	108.39	*	-
	NT2RM2000973	104.1	257.68	179.14	123.46	180.6	103.68	80.15	78.99	82.31		
	NT2RM2000983	44.05	48.21	49.47	28.35	25.15	19.95	21.95	14.49	14.64	** **	- -
	NT2RM2000984	14.04	16.71	14.65	9.74	5.64	4.75	5.34	3.15	3.97	** **	- -
45	NT2RM2000994	36.03	29.02	31.03	22.45	13.36	11.64	9.34	22.92	20.85	* *	- -
	NT2RM2001004	87.18	99.34	120.85	105.48	104.79	118.66	75.52	78.54	88.2		
	NT2RM2001022	465.07	829.81	488.65	279.37	461.13	292.23	411.29	508	337.91		
	NT2RM2001035	119.48	128.64	135.94	69.22	98.88	76.14	85.64	92.65	116	* *	- -
	NT2RM2001038	30.1	53.09	55.15	35.29	31.06	27.29	33.8	26.97	56.89		
	NT2RM2001043	17.78	17.9	22.19	14.07	14.29	12.96	14.08	13.2	13.27	* *	- -
50	NT2RM2001050	16.36	10.75	11.34	9.74	5.11	3.09	5.76	4.31	3.75	*	-
	NT2RM2001055	7.49	7.2	8.19	5.79	2.69	1.59	3.42	2.92	3.04	* **	- -
	NT2RM2001065	18.94	19.43	21.19	16.28	10.87	7.49	3.44	9.8	10.97	* **	- -
	NT2RM2001075	352.66	309.42	355.51	314.53	188.83	298.2	178.61	439.36	452.62		
	NT2RM2001083	7.08	9.84	12.58	9.39	11.2	5.35	6.46	10.07	4.34		
	NT2RM2001100	132.53	127.49	140.57	168.49	157.74	186.39	108.36	105.08	125.06	* *	+ -
55	NT2RM2001105	19.7	23.88	18.02	9.87	6.9	7.5	7.31	7.33	7.99	** **	- -
	NT2RM2001109	20.13	43.33	44.99	10.64	14.91	10.34	19.4	14.81	19.72	*	-
	NT2RM2001110	32.89	28.27	36.47	39.19	23.1	32.21	27.03	23.61	20.38	*	-

Table 406

	NT2RM1001103	12.11	14.64	19.71	7.84	12.92	9.54	14.04	7.04	13.96			
	NT2RM1001105	4.22	7.99	12.4	1.52	4.66	3.36	1.61	1.94	3.34			
5	NT2RM1001112	5.26	6.44	10.76	4.28	9.37	3.52	5.38	1.59	3.46			
	NT2RM1001115	7.42	6.2	16.09	7.99	9.99	7.63	6.92	3.96	3.94			
	NT2RM1001122	13.95	16.42	25.8	14.65	15.41	13.67	7.73	7.37	13.09			
	NT2RM1001136	4.38	9.14	8.72	2.16	2.27	3.09	2.35	0.89	0.34	*	*	-
	NT2RM1001139	10.44	16.1	18.58	8.95	9.3	8.97	8.52	6.5	7.34	*	*	-
10	NT2RM2000003	3.68	3.81	8.36	4.11	1.99	1.48	3.39	1.06	1.76			
	NT2RM2000006	19.96	26.87	32.42	17.36	19.88	13.64	34.18	15.67	16.17			
	NT2RM2000010	13.11	32.27	43.36	20.14	26.86	16.98	10.37	4.84	2.67			
	NT2RM2000013	111.85	65.82	177.02	150.21	118.48	155.88	73.81	57	69.05			
	NT2RM2000030	47.5	43.12	78.78	78.26	63.56	52.76	55.62	28.2	43.38			
	NT2RM2000032	27.93	34.32	46.51	34.09	25.35	35.27	22.38	22.42	18.15			
15	NT2RM2000039	15.44	18.82	11.91	10.78	6.19	11.58	11	10.28	4.44			
	NT2RM2000042	22.91	29.95	35.12	39.39	23.39	35.58	21.42	15.87	21.82			
	NT2RM2000082	3.58	4.4	7.65	2.5	2.61	1.79	2.27	5.17	2.19			
	NT2RM2000093	9.7	29.83	14.41	14.47	18.41	7.29	12.45	10.38	11.19			
	NT2RM2000101	92.12	99.66	118.62	114.91	128.83	118.95	69.62	52.13	78.59	*		-
	NT2RM2000104	62.95	69.41	129.33	110.95	90.54	74.09	72.52	52.94	63.78			
20	NT2RM2000124	8.36	13.36	22.64	10.6	20.3	10.85	19.19	7.38	15.25			
	NT2RM2000155	8.61	19.1	14.45	7.85	7.12	5.48	4.06	1.65	4.16	*		-
	NT2RM2000191	45.98	51.15	77.89	79.82	71.61	59.62	35.22	25.83	44.53			
	NT2RM2000192	4.72	2.97	6.83	2.91	2.72	2.95	2.76	1.61	1.47			
	NT2RM2000239	13.1	14.91	17.36	3.56	5.21	3.86	9.92	3.36	2.39	**	*	-
	NT2RM2000241	10.79	19.6	19.09	8.89	13.98	6.49	11.46	3.01	8.51			
25	NT2RM2000250	5.15	5.11	16.92	5.37	6.41	3.58	6.45	1.4	5.59			
	NT2RM2000259	7.84	17.03	22.56	4.92	6.83	6.63	8.62	2.03	2.42			
	NT2RM2000260	9.47	10.65	9.59	4.97	5.31	6.44	6.43	4.5	4.7	**	**	-
	NT2RM2000265	1.13	1.27	5.15	0.34	1.13	1.33	1.14	0	0.82			
	NT2RM2000287	47.66	55.1	96.28	31.39	36.17	31.93	36.1	33.17	17.11			
	NT2RM2000306	66.48	31.19	72.95	23.2	54.18	20.2	25.43	22.47	8.61			
30	NT2RM2000312	32.44	17.61	49.51	8.46	13.33	15.12	25.48	21.27	10.02			
	NT2RM2000322	7.99	6.33	16.54	4.19	5.04	3.2	3.47	0.77	1.48			
	NT2RM2000343	85.23	93.06	177.62	122.74	138.14	108.3	83.09	46.31	57.15			
	NT2RM2000359	6.22	10.59	23.8	9.56	9.62	5.13	12.03	2.74	6.22			
	NT2RM2000362	121.28	140.62	246.7	173.62	223.63	249.97	108	78.11	121.29			
	NT2RM2000363	3.84	6.52	9.96	6.57	3.72	4.83	5.89	2.02	2.82			
35	NT2RM2000368	26.44	19.81	43.02	11.49	11.34	10.46	8.57	3.39	3.24	*	*	-
	NT2RM2000371	547.25	508.14	752.99	148.13	425.69	273.34	355.28	228.93	135.68	*	*	-
	NT2RM2000374	12.96	13.63	33.5	14.47	12.05	8.49	18.75	11.86	10.92			
	NT2RM2000387	30.78	60.24	91.33	10.11	11.64	24.36	10.29	9.16	3.11	*		-
	NT2RM2000393	10.16	9.5	18.93	3.77	7.23	5.64	6.78	3.44	2.74			
	NT2RM2000395	3.42	4.7	7.68	4.26	7.48	3.65	6.98	2.09	2.77			
40	NT2RM2000402	19.66	22.97	34.41	26.49	10.93	25.33	8.72	3.41	2.72	*		-
	NT2RM2000405	3.7	3.95	7.31	4.5	3.17	3.25	3.78	1.94	0.68			
	NT2RM2000407	6.18	7.05	11.03	8.23	7.83	7.22	4.36	3.58	3.42	*		-
	NT2RM2000410	6.69	5.24	14.77	10.81	8.59	8.57	4.92	7	5.39			
	NT2RM2000420	12.46	12.68	20.53	7.85	6.53	12.66	7.69	6.33	9.37			
	NT2RM2000422	13.42	17.5	14.12	8.88	9.58	15.24	10	8.21	5.8	*		-
45	NT2RM2000423	15.54	22.01	50.31	29.73	23.33	19.31	31.02	12.98	24.15			
	NT2RM2000452	3.39	4.23	12.53	5.47	7.75	4.42	5.91	1.75	4.57			
	NT2RM2000469	3.15	2.51	7.63	7.05	2.6	1.83	2.99	3.89	3.38			
	NT2RM2000480	12.63	13.33	17.91	18.49	9.81	12.7	9.63	10.71	10.64			
	NT2RM2000487	6.31	3.98	9.54	8.32	2.31	5.44	4.36	6.15	6.13			
	NT2RM2000502	7.65	10.01	17.15	15.2	7.83	12.64	18.37	19.29	16.37			
50	NT2RM2000504	8.62	13.61	12.8	14.35	8.61	11.65	11.39	5.05	12.11			
	NT2RM2000514	3.24	4.13	6.05	5.11	2.16	2.21	3.1	2.79	1.66			
	NT2RM2000522	0.79	1.87	2.28	5.66	1.51	1.86	2.44	0.56	1.28			
	NT2RM2000540	36.37	16.56	24.89	25.09	10.1	17.62	15.11	17.92	16.53			
	NT2RM2000555	10.96	7.91	19.23	7.94	6.73	6.21	4.49	7.85	6.72			
	NT2RM2000566	35.88	29.95	48.96	40.83	31.02	33.77	42.8	32.01	34.75			
55	NT2RM2000567	5.01	5.28	9.14	7.2	2.55	2.31	2.41	4.02	2.26			
	NT2RM2000589	11.75	12.7	16.59	10.42	10.16	8.54	12.04	11.54	10.63			
	NT2RM2000577	23.09	36.16	46.55	20.42	16.61	17.63	9.65	9.61	18.29	*		-

Table 405

	NT2RM1000800	8.8	11.31	17.21	5.12	7.9	6.45	6.62	2.19	3.3	*	-
	NT2RM1000802	37.09	50.36	74.58	36.6	34.84	28.86	14.15	4.56	9.84	*	-
5	NT2RM1000811	12.36	9.86	24.52	8.71	10.02	9.58	14.12	5.95	8.7		
	NT2RM1000826	30.88	47.18	81.3	44.32	50.77	35.09	43.27	14.54	16.43		
	NT2RM1000829	19.85	28.15	42.49	15.46	18.32	22.11	18.5	10.73	22.04		
	NT2RM1000831	641.62	637.04	913.82	275.83	388.75	407.08	509.83	541.98	257.92	*	-
	NT2RM1000833	131.01	121.94	243.56	117.55	172.46	143.86	179.55	110.44	138.22		
	NT2RM1000834	26.33	24.31	2.89	0	0.65	2.02	26.66	20.39	19.97		
10	NT2RM1000841	24.25	46.24	73.12	11.51	14.12	14.5	18.59	16.34	8.01		
	NT2RM1000848	14.13	26.98	37.82	13	21.43	18.99	20.71	11.34	13.28		
	NT2RM1000850	24.68	26.73	36.64	21.15	32.91	25.01	46.02	20.2	29.9		
	NT2RM1000852	13.97	15.95	23.78	18.75	27.94	15.23	31.21	18.1	21.46		
	NT2RM1000853	4.2	3.72	6.66	7.37	4.53	5.05	3.89	2.23	2.94		
	NT2RM1000855	68.98	71.69	82.2	36.85	37.1	51.79	74.89	42.55	21.65	**	-
15	NT2RM1000857	18.68	28.42	44.59	21.2	21.59	23.12	20.54	14.01	19.44	*	-
	NT2RM1000858	19.44	18.6	24.07	17.38	13.26	16.79	12.93	7.9	10.41	*	-
	NT2RM1000867	47.06	39.97	81.53	51.39	22.67	30.69	24.25	41.66	28.02		
	NT2RM1000874	15.53	21.9	42.18	11.95	15.55	12.17	9.14	4.05	6.39		
	NT2RM1000882	8.95	14.01	25.71	7.66	9.72	9.63	17.06	6.72	11.92		
20	NT2RM1000883	14.26	11.55	31.41	10.17	21.02	6.37	28.25	10.83	25.46	**	-
	NT2RM1000885	24.96	25.3	31.55	8.98	10.57	16.46	9.36	11.69	11.59	**	-
	NT2RM1000893	28.28	40.4	46.33	35.83	16.57	27.75	26.99	25.66	21.62		
	NT2RM1000894	11.16	14.78	12.12	9.67	5.22	9.03	7.62	4.24	5.7	**	-
	NT2RM1000898	16.49	18.47	21.09	20.98	15.98	15.23	9.37	6.69	9.26	**	-
	NT2RM1000899	5.15	4.63	7.57	5.14	7.26	5.6	4.73	2.38	2.79		
25	NT2RM1000905	294.53	321.81	294.85	264.7	213.45	208.71	204.71	142.52	169.18	*	-
	NT2RM1000910	73.69	67.82	69.43	87.95	125.51	72.31	61.51	49.51	44.37	*	-
	NT2RM1000914	51.65	67.13	49.73	46.16	55.8	39.43	58.15	36.24	73.81		
	NT2RM1000919	31.63	34.55	49.42	11.97	21.16	18.78	16.17	26.42	28.42	*	-
	NT2RM1000921	6.87	9.33	11.95	7.94	8.86	8.21	8.17	4.59	6.66		
	NT2RM1000922	13.29	8.81	21.03	4.57	14.42	11.98	8.29	7.63	6.27		
30	NT2RM1000924	4.71	2.79	7.57	4.61	10.85	2.89	2.79	1.75	0.64		
	NT2RM1000927	6.24	8.21	11.68	5.06	6.28	6.3	9.48	6.11	7.76		
	NT2RM1000951	43.38	31.87	51.83	47.29	42.21	49.1	23.28	12.15	25.07	*	-
	NT2RM1000956	14.05	17.39	13.26	12.15	19.17	13.5	7.97	3.02	6.22	**	-
	NT2RM1000960	27.19	36.09	41.59	49.18	53.65	39.11	40.71	16.46	20.8		
	NT2RM1000961	31.7	37.01	55	10.53	21.75	21.52	13.1	26.97	26.19	*	-
35	NT2RM1000962	27.85	38.63	33.9	11.13	12.72	11.99	4.23	8.24	8.72	**	-
	NT2RM1000973	43.5	24.74	82.26	40.71	70.49	56.15	24.22	34.19	11.88		
	NT2RM1000978	0.68	0.64	4.81	0	0.42	0.84	0.61	2.06	0.63		
	NT2RM1000982	16.77	17.99	20.27	10.66	18.58	11.05	7.48	6.36	6.54	**	-
	NT2RM1000991	7.55	12.77	12.19	10.81	11.04	11.17	5.79	6.02	8.13		
	NT2RM1000994	39.31	47	23.95	9.37	52.07	40.12	12.98	9.18	11.27	*	-
40	NT2RM1001002	32.74	42.78	50.1	43.61	48.24	33.17	37.45	15.22	13.9		
	NT2RM1001003	22.14	23.85	35.08	16.55	19.31	27.31	8.71	13.75	14.54	*	-
	NT2RM1001008	2.98	2.17	4.26	1.07	1.49	2.65	1.82	2.32	1.41		
	NT2RM1001011	24.49	19.42	39.5	9.16	16.59	21.83	7.1	5.35	13.23	*	-
	NT2RM1001013	7.58	6.4	15.54	3.26	7.12	6.38	8.21	10.22	6.35		
	NT2RM1001017	2.99	2.36	7.63	0.68	1.65	2.36	1.32	0	2.89		
45	NT2RM1001018	389.38	678.69	541.36	131	345.59	266.19	208.53	225.12	141.21	*	-
	NT2RM1001026	7.53	12.1	18.29	9.07	8.15	5.48	3.88	1.52	2.01	*	-
	NT2RM1001028	4.9	7.17	9.41	8.47	12.06	5.7	11.7	4.86	5.5		
	NT2RM1001043	26.98	28.41	30.47	32.28	25.93	18.1	7.33	9.19	17.19	**	-
	NT2RM1001044	13.61	15.79	12.73	5.11	11.56	9.29	6.83	6	8.45	**	-
	NT2RM1001059	17.62	17.16	21.39	29.41	18.53	17.53	21.59	10.53	22.05		
50	NT2RM1001063	4.42	2.79	10.14	1.51	2.68	1.95	3.55	2.61	2.37		
	NT2RM1001066	5.44	6.05	11.22	3.97	4.37	2.62	5.3	2.25	4.68		
	NT2RM1001072	2.1	2.09	9.94	0.59	2.67	1.88	2.14	0.14	1.84		
	NT2RM1001074	4.98	10.47	11.14	5.42	6.47	3.23	4.47	1.73	1.4	*	-
	NT2RM1001076	7.28	10.41	10.61	10.91	16.92	6.26	16.73	6.71	6.46		
	NT2RM1001082	15.79	21.18	26.04	16.26	13.72	10.05	11.58	9.75	14.56		
55	NT2RM1001085	1.82	1.24	5.39	0.21	0.47	1.19	1.3	0.55	1.17		
	NT2RM1001092	38.97	56.81	56.39	54.52	49.61	49.24	59.76	48.35	50.33		
	NT2RM1001102	10.81	9.79	17.74	9.01	8	8.42	8.99	8.26	5.92		

Table 404

	NT2RM1000354	0.4	1.68	2.48	0.51	1.02	0.95	0.87	0.21	0.21			
	NT2RM1000355	18.03	21.12	26.82	24.57	24.38	11.58	16.5	8.01	13.65			
5	NT2RM1000361	2.57	1.81	2.89	1.57	3.04	3.2	2.49	0.23	2.67			
	NT2RM1000365	0.53	2.31	3.79	0	3.39	1.9	2.3	0	1			
	NT2RM1000372	55.88	63.08	56.77	80.59	90.54	75.31	135.63	66.87	19.91	**	+	
	NT2RM1000377	29.22	32.95	43.33	46.89	24.42	30.81	18.03	18.61	14.45	*	-	
	NT2RM1000388	3.72	3.27	4.82	4.15	1.73	3.91	2.28	2.59	2.31	*	-	
10	NT2RM1000394	1.56	1.31	3.56	1.15	0.93	0.79	1.9	2.42	1.43			
	NT2RM1000399	5.42	3.53	8.67	4.2	3.61	3.12	6.84	2.24	2.92			
	NT2RM1000407	1.68	1.87	7.64	2.01	1.44	1.69	2.06	1.13	3.04			
	NT2RM1000421	0.74	0.94	3.14	1.35	1.14	2	1.61	0	0.06			
	NT2RM1000422	134.24	176.38	156.33	143.97	232.16	154.32	116.71	110.46	75.99	*	-	
	NT2RM1000430	4.06	4.21	6.87	5.11	7.78	3.82	6.49	1.64	1.64			
15	NT2RM1000462	17.49	21.49	23.94	21.57	15.84	20.24	13.6	12.18	18.48			
	NT2RM1000489	11.77	14.5	12.66	4.08	5.8	6.16	2.2	2.63	4.58	**	**	-
	NT2RM1000512	50.79	65.54	89.54	15.65	74.29	59.09	19.69	52.23	56.59			
	NT2RM1000519	108.54	108.15	106.36	60.23	134.25	94.69	58.45	58.56	60.96	**	-	
	NT2RM1000527	51.37	48.34	68.1	100.86	64	65.36	54.81	19.74	44.25			
	NT2RM1000539	19.83	27.84	33.74	35.02	28.56	31.96	16.52	8.8	16.69			
20	NT2RM1000542	2.91	2.41	11.46	0.76	1.72	0.95	1.66	0.17	0.24			
	NT2RM1000553	49.27	35.57	63.05	73.97	60.28	58.45	40.5	24.9	19.69			
	NT2RM1000555	85.13	121.39	124.03	87.28	76.76	74.83	41.33	57.49	77.7	*	-	
	NT2RM1000558	26.75	25.54	23.49	36.55	31.73	36.63	21.2	19.34	35.09	**	+	
	NT2RM1000563	9.05	10.86	14.09	10.89	9.42	9.24	7.48	6.52	10.32			
	NT2RM1000566	6.73	8.38	15.32	6.55	8.24	9.4	8.98	6.08	9.24			
25	NT2RM1000570	125.7	201.18	105.25	41.82	195.51	106.3	185	132.56	153.14			
	NT2RM1000571	50.08	74.69	65.76	14.46	84.43	39.98	43.45	24.14	6.89	*	-	
	NT2RM1000574	2.53	1.65	9.83	1.89	3.52	1.98	1.99	0.19	1.58			
	NT2RM1000580	7.86	7.46	8.12	4.52	12.24	4.74	8.38	3.22	4.57			
	NT2RM1000620	22.3	25.58	42.74	35.73	32.55	28.94	25.27	21.57	27.34			
	NT2RM1000623	1.13	1.82	2.51	0.54	0.62	1.17	0.35	0.18	1.57			
30	NT2RM1000630	3.11	2.76	5.89	2.21	1.76	2.12	2.49	0.75	1.9			
	NT2RM1000633	91.32	89.94	117.97	56.97	108.85	75.65	98.98	107.05	51.97			
	NT2RM1000634	11.16	13.34	24.68	9.79	14.73	9.21	12.08	6.93	12.7			
	NT2RM1000642	12.72	15.83	20.12	6.07	8.95	6.42	3.63	2.38	4.98	*	**	-
	NT2RM1000647	58.32	165.39	134.62	67.85	112.06	61.29	71.96	52.24	87.23			
	NT2RM1000648	7.26	13.24	8.89	5.57	12.68	6.6	14.52	3.49	7.15			
35	NT2RM1000650	6.62	9.44	9.46	2.83	6.47	6.87	3.69	5.63	4.1	*	-	
	NT2RM1000661	15.88	18.26	26.1	21.53	19.01	24.34	8.08	7.55	8.54	*	-	
	NT2RM1000666	1.96	1.18	6.92	1.86	1.52	1.45	0.88	0.27	0.92			
	NT2RM1000669	17.26	18.78	20.02	4.09	4.91	3.69	2.13	1.18	2.17	**	**	-
	NT2RM1000672	13.67	18.24	42.69	19.37	17.89	17.56	7.45	2.82	5.99			
	NT2RM1000681	9.08	16.58	18.31	9.41	17.68	12.92	13.31	5.99	9.03			
40	NT2RM1000691	4.68	7.55	17.93	4	7.1	7.03	5.99	3.47	5.58			
	NT2RM1000698	12.42	13.98	20.53	4.92	32.5	5.23	15.99	3.16	6.38			
	NT2RM1000699	5.67	7.09	9.09	7.34	5.78	7.37	2.45	1.86	3.56	*	-	
	NT2RM1000702	25.96	30.69	39.69	30.16	23.23	32.77	24.6	18.35	22.21			
	NT2RM1000703	32.53	88.64	53.23	36.64	56.43	43.04	42.73	50.17	17.73			
	NT2RM1000704	43.17	84	45.95	47.88	42.53	19.03	39.69	25.24	10.87			
45	NT2RM1000726	5.67	6.22	17.4	3.29	4.23	3.21	4.45	0.93	1.78			
	NT2RM1000731	21.17	21.67	40.88	23.29	32.82	14.51	24.97	11.69	10.31			
	NT2RM1000741	3.88	5.18	10.63	4.4	7.39	2.14	6.17	0	1.39			
	NT2RM1000742	28.88	28.51	36.7	38.75	29.44	32.74	34.48	36.03	27.02			
	NT2RM1000744	18.57	21.62	27.17	23.44	20.36	17.29	20.61	18.9	23.17			
	NT2RM1000746	7.85	8.11	11.72	4.17	4.62	5.27	5.53	3.03	1.14	*	*	-
50	NT2RM1000747	30.16	38.37	65.95	28.71	33.95	34.03	30.94	18.08	23.76			
	NT2RM1000752	6.43	11.58	14.15	3.61	7.68	5.76	4.96	2.69	4.73			
	NT2RM1000767	82.05	87.93	86.38	98.35	104.39	73.49	112.1	59.62	78.42			
	NT2RM1000770	9.08	11.59	19.18	5.23	6.92	2.95	3.61	2.19	4.43	*	-	
	NT2RM1000772	0.81	1.85	7.5	1.21	3.6	1.86	2.53	0	2.7			
	NT2RM1000779	107.98	131.01	176.51	63.61	94.24	109.72	76.84	69.35	114.65			
55	NT2RM1000780	6.52	7.85	9.21	2.66	6.05	5.1	2.74	1.38	1.99	**	-	
	NT2RM1000781	6.79	4.22	9.28	1.5	2.8	0.61	2.07	1.26	0.75	*	*	-
	NT2RM1000789	63.83	77.27	95.07	93.46	102.47	101.3	60.5	47.17	81.09			

Table 403

5	MAHMA1003157	9.8	11.92	16.49	8.35	7.14	7	5.32	5.07	5.53	*	-
	MAHMA1003163	6.69	5.07	11.11	7.07	6.08	5.54	2.67	4.3	4.05	*	-
	MAHMA1003164	9.65	8.42	12.05	4.3	5.95	3.77	2.65	1.16	3.38	*	-
	MAHMA1003166	13.16	17.5	21.96	14.35	15.87	10.73	8.74	5.62	7.88	*	-
	NB9N31000010	22.17	30.34	35	8.06	26.76	14.32	17.42	5.42	10.61	*	-
10	NB9N31000016	6.11	10.91	22.1	11.04	8.39	6.1	4.87	1.93	11.92	*	-
	NB9N31000043	10.01	15.51	20.02	13.55	11.04	7.12	6.17	3.99	7.07	*	-
	NB9N31000045	170.84	205.1	142.2	172.26	138.41	173.74	84.94	88.64	18.56	*	-
	NB9N31000054	36.72	46.14	51.85	45.82	42.67	36.97	50.28	31.68	29.41	*	-
	NB9N31000076	7.95	11.11	16	8.53	7.69	7.72	11.59	4.88	6.85	*	-
15	NB9N31000086	4.15	2.68	11.91	2.36	4	2.05	3.15	0.81	2.03	*	-
	NT2RM1000001	7.05	9.84	23.41	7.46	11.26	6.73	9.81	4.19	5.49	*	-
	NT2RM1000018	6.2	11.24	12.18	4.32	8.37	5.02	4.96	1.12	2.49	*	-
	NT2RM1000032	6.17	20.65	14.02	3.2	3.21	6.55	7.1	3.44	6.93	*	-
	NT2RM1000035	5.32	12.32	11.65	5.42	4.73	4.54	3.82	4.31	2.73	*	-
20	NT2RM1000037	6.94	9.49	8.81	3.59	2.53	4.32	3.57	2.25	1.49	**	-
	NT2RM1000039	17.35	20.78	18.95	11.31	10.08	7.38	8.83	5.7	8.35	**	-
	NT2RM1000042	285.1	504.24	359.39	438.78	386.15	125.88	306.06	229.38	295.8	**	-
	NT2RM1000055	3.2	5.45	10.27	2.35	2.5	0.71	3.16	1.31	1.79	*	-
	NT2RM1000059	6.89	5.26	12.3	2.6	4.12	1.14	3.74	1.4	3.81	*	-
25	NT2RM1000062	3.25	17.72	14.57	2.77	4.02	2.38	6.97	0.66	3.27	*	-
	NT2RM1000065	129.93	146.35	187.39	51.11	120.84	109.9	91.66	120.6	204.01	*	-
	NT2RM1000066	8.15	10.03	11.93	3.29	5.25	5.44	3.51	4.17	3.47	*	-
	NT2RM1000071	663.12	662.77	828.24	110.41	580.78	259.28	351.73	417.5	139.33	*	-
	NT2RM1000080	24.34	23.12	24.37	3.56	11.29	8.21	9.27	5.24	4.03	**	-
30	NT2RM1000086	10.86	12.97	11.53	2.83	5.32	3.9	5.64	4.4	2.88	**	-
	NT2RM1000092	20.13	29.95	35.15	36.08	29.43	24.61	19.3	14.65	20.49	*	-
	NT2RM1000118	1.37	1.52	9.04	0	0.67	0	0.36	0	0.79	*	-
	NT2RM1000119	1.84	9.53	13.52	3.77	3.18	2.59	3.62	0	1.09	*	-
	NT2RM1000121	4.53	3.89	4.67	2.15	3.4	5.49	1.08	0.78	1.77	**	-
35	NT2RM1000122	12.06	9.02	14.76	4.72	5.56	4.77	3.17	7.39	6.45	*	-
	NT2RM1000127	4.86	3.09	8.85	2.65	2.99	2.48	1.52	1.46	1.57	*	-
	NT2RM1000131	2.61	0.69	3.44	0.91	1.56	0.43	1.25	0.9	1.02	*	-
	NT2RM1000132	14.01	8.86	18.39	14.15	9.4	7.78	17.49	15	6.09	*	-
	NT2RM1000153	5.01	3.78	14.31	1.42	4.08	2.81	5.68	1.59	2.53	*	-
40	NT2RM1000184	281.4	361.08	212.97	228.94	161.22	121.63	212.54	317.63	41.9	*	-
	NT2RM1000186	2.47	4.7	12.56	1.93	2.64	2.03	5.08	0.92	2.79	*	-
	NT2RM1000187	13.51	16.82	27.58	4.35	7.1	11.13	3.58	3.22	6.78	*	-
	NT2RM1000199	2.37	2.51	5.33	3.1	3.18	3.27	2.76	0.66	1.97	*	-
	NT2RM1000213	14.38	13.17	27	9.3	8.17	9.61	8.4	3.99	3.63	*	-
45	NT2RM1000215	127.65	93.55	105.56	47	125.55	80	93.27	92.32	37.71	*	-
	NT2RM1000218	17.5	12.81	37.21	8.7	7.66	13.12	12.93	9.58	4.51	*	-
	NT2RM1000224	38.8	21.38	99.03	21.96	23.06	30.11	56.22	41.03	10.49	*	-
	NT2RM1000236	128.23	106.27	224.67	36.44	158.46	117.76	233.25	180.75	49.09	*	-
	NT2RM1000242	0.21	0.74	2.56	0	1.12	0	0.56	0	0.76	*	-
50	NT2RM1000244	7.07	7.69	14.14	11.09	6.96	9.63	3.01	2.46	3.87	*	-
	NT2RM1000252	3.63	3.86	6.34	5.75	3.24	4.11	5.02	2.57	2.44	*	-
	NT2RM1000256	13.89	13.11	17.86	12.49	12.19	15.55	9.96	9.49	3.8	*	-
	NT2RM1000257	7.26	4.79	22.28	12.27	7.82	12.93	3.88	4.6	1.41	*	-
	NT2RM1000260	44.37	50.35	69.11	33.91	47.16	50.17	26.8	25.27	25.69	*	-
55	NT2RM1000269	4.48	3.66	6.5	3.35	3.23	4.63	2.09	1.54	1.92	*	-
	NT2RM1000271	2.07	1.72	5.77	1.75	2.06	0.52	0.83	0.09	0.38	*	-
	NT2RM1000272	93.8	125.17	184.07	85.39	186.36	110.41	166.71	67.77	97.63	*	-
	NT2RM1000273	44.49	55.95	63.71	32.07	31.24	23.66	15.32	26.55	33.94	*	-
	NT2RM1000274	455.63	719.55	944.44	916.81	390.33	402.82	570.68	746.39	848.53	*	-
55	NT2RM1000280	6.37	12.22	16.78	15.15	6.91	15.12	10.36	3.5	4.59	*	-
	NT2RM1000295	0.98	1.35	3.23	1.46	0.9	2.46	0.9	0.93	3.62	*	-
	NT2RM1000300	6.74	6.95	12.1	10.61	8.44	10.7	5.89	3.63	7.13	*	-
	NT2RM1000304	737.06	1023.4	718.47	1038.7	1097.8	469.75	946.67	780.24	888.74	*	-
	NT2RM1000314	4.57	4.01	6.01	3.26	3.88	4.08	4.92	1.32	3.56	*	-
55	NT2RM1000318	421.37	294.99	289.4	315.7	574.37	386.7	448.45	209.13	302.89	*	-
	NT2RM1000335	11.96	11.64	18.91	9.64	8.09	12	6.06	5.29	8.48	*	-
	NT2RM1000341	1.94	1.67	2.31	1.53	0.88	0.82	1.64	0.39	2.21	*	-
	NT2RM1000350	8.91	10.88	13.15	11.12	8.83	9.83	7.57	4.14	7.11	*	-

Table 402

	MAMMA1002890	11.31	9.48	18.16	8.56	8.49	19.82	9.15	5.95	8.48		
	MAMMA1002892	13.36	11.62	14.89	12.15	11.02	9.05	7.92	7.62	11.8		
5	MAMMA1002893	8.64	14.35	13.28	7.89	13.97	5.96	10.13	7.23	1.86		
	MAMMA1002895	3.34	5.32	14.4	3.46	5.81	1.87	4.28	2.34	3.5		
	MAMMA1002898	4.08	5.01	17.31	2.16	5.01	1.78	3.17	0.38	2.02		
	MAMMA1002905	4	11.4	20	7.75	8.82	9.71	6.71	1.7	4.46		
	MAMMA1002906	2.68	1.92	5.05	4.09	2.8	3.58	1.97	0.54	1.64		
10	MAMMA1002908	14.98	16.18	26.75	12.12	15.62	14.31	16.05	8.69	18.72		
	MAMMA1002909	30.34	34.63	68.19	28.69	18.89	22.65	34.89	20.01	32.36		
	MAMMA1002918	13	10.64	27.29	16.01	18.07	12.45	12.7	8.75	10.42		
	MAMMA1002925	49.1	23.96	62.42	27.59	36.92	32.28	67.91	43.67	13.09		
	MAMMA1002926	25.41	28.93	57.26	38.61	37.53	35.54	28.57	14.19	28.89		
	MAMMA1002930	27	34.75	73.56	25.58	25.59	23.42	47.23	26.24	37.21		
15	MAMMA1002937	65.34	59.27	78.68	106.43	76.67	71.14	57.91	26.1	50.41		
	MAMMA1002938	1.94	2.84	7.68	5.03	4.18	4.4	2.05	0.91	2.65		
	MAMMA1002941	7.09	6.07	17.41	12.34	8.56	8.04	6.56	3.92	5.66		
	MAMMA1002947	16.06	20.72	33.43	32.88	25.81	27	19.39	13.57	20.15		
	MAMMA1002964	14.52	21.96	48.58	29.27	22.13	24.66	18.71	13.88	18.49		
	MAMMA1002967	6.96	4.19	15.96	5.47	7.15	5.12	3.82	5.07	4.36		
20	MAMMA1002970	16.09	29.77	65.22	27.91	25.08	19.79	28.99	12.64	23.95		
	MAMMA1002971	4.02	4.7	8.35	3.51	4.7	2.27	5.82	2.89	2.46		
	MAMMA1002972	3.94	3.63	9.75	6.43	4.91	5.11	4.27	0.9	3.28		
	MAMMA1002973	20.96	22.17	31	24.13	16.86	22.18	16.66	11.21	13.94	*	-
	MAMMA1002979	129.92	194.32	219.31	151.08	71.59	113.43	55.53	74.31	23.22	*	-
	MAMMA1002982	2.18	1.25	7.22	2.29	3.22	3.34	5.33	0.9	3.23		
25	MAMMA1002987	10.35	12.2	15.88	9.6	9.29	9.6	13.48	8.42	11.93		
	MAMMA1003003	10.92	12.48	21.55	9.72	10.82	10.86	12.69	9.52	10.22		
	MAMMA1003004	9.35	12	11.41	6.36	11.7	10.34	5.34	4.84	8.09	*	-
	MAMMA1003007	5.35	7.51	8.39	5.22	6.88	6.35	8.36	3.78	6.7		
	MAMMA1003011	3.33	3.62	5.57	4.14	2.91	4.01	2.44	1.93	2.62		
	MAMMA1003013	52.58	50.01	73.47	127.57	69.44	113.36	42.21	37.2	47.45		
30	MAMMA1003015	4.4	9.22	10.22	6.68	9.17	6.66	6.79	4.74	5.08		
	MAMMA1003019	6	3.52	10.72	6.3	3.92	5.12	4.27	1.82	2.44		
	MAMMA1003020	9.29	17.63	21.88	11.76	13.24	12.14	9.04	4.99	7.31		
	MAMMA1003026	2.19	2.41	7.93	1.47	3.27	2.09	1.98	0.75	2.98		
	MAMMA1003031	12.83	17.1	23.06	9.93	17.27	14.35	13.73	7.28	15.8		
	MAMMA1003033	7.59	11.58	13.68	6.87	12.75	8.45	8.32	3.56	9.08		
35	MAMMA1003035	14.44	18.88	35.58	17.37	11.86	13.06	6.41	9.44	10.62		
	MAMMA1003039	5.57	9.4	15.11	7.17	7.63	9.67	4.3	4.16	6.95		
	MAMMA1003040	22.42	25.75	39.61	22.72	19.76	23.47	30.8	25.19	32.64		
	MAMMA1003044	23.58	26.13	43.98	21.92	19.68	21.85	23.01	22.17	23.66		
	MAMMA1003047	25.89	30.87	33.12	18.34	23.66	19.64	30.18	18.74	28.8	*	-
	MAMMA1003049	2.33	2.17	8.04	1.08	1.14	1.06	1.63	0.08	1.3		
40	MAMMA1003055	17.22	25.14	35.08	23.45	17.28	15.29	23.26	16.76	22.03		
	MAMMA1003056	1.24	3.89	8.92	1.85	1.6	1.77	4.37	0	1.6		
	MAMMA1003057	7.01	11.35	14.1	7.31	6.88	7.83	3.86	5.43	8.7		
	MAMMA1003066	16.42	21.28	30.29	14.08	16.97	16.54	12.01	11.17	25.5		
	MAMMA1003075	6.07	7.85	12.07	6.85	6.1	6.14	7.15	5.26	7.72		
	MAMMA1003089	37.69	44.94	70.13	31.56	35.51	31.8	38.3	22.85	38.78		
45	MAMMA1003092	2.7	6.77	9.29	2.59	2.39	2.86	1.65	0.84	2.49		
	MAMMA1003095	11.71	16.36	22.28	5.96	7.13	4.55	3.24	2.25	2.45	*	*
	MAMMA1003099	12.83	14.7	25.29	10.89	15.86	9.22	14.89	11.45	12.15		
	MAMMA1003102	3.77	5.08	9.09	1.68	4.54	2.6	2.97	0.85	1.92		
	MAMMA1003104	2.23	3.23	4.66	5.3	4.9	4.18	1.01	1.61	4.75		
	MAMMA1003113	12.38	25.98	27.12	11.65	17.89	10.18	7.19	8.58	13.27		
50	MAMMA1003126	20.48	25.06	25.6	15.4	13.83	15.96	10.44	8.97	17.98	**	*
	MAMMA1003127	11.67	12.89	22.02	11.38	12.69	10.66	9.71	8.8	11.4		
	MAMMA1003131	8.9	14.73	17.42	6.88	7.18	8.51	7.06	3.18	4.59	*	-
	MAMMA1003135	3.06	2.9	10.48	1.24	2.69	1.3	3	1.87	1.55		
	MAMMA1003140	5.46	6.32	15.55	2.89	4.63	3.63	3.24	3.7	5.1		
	MAMMA1003146	4.31	6.5	9.12	2.65	6.5	3.58	4.41	3.82	3.28		
55	MAMMA1003150	4.32	7.47	10.56	5.78	4.99	3.76	3.2	3.41	5.87		
	MAMMA1003154	2.93	3.77	5.99	2.37	2.94	2.85	2.68	0.86	2.1		
	MAMMA1003155	35.26	33.7	28.41	18.29	14.13	13.99	27.83	12.47	20.55	**	-

Table 401

	MAMMA1002655	11.4	14.99	17.9	15.12	13.04	14.83	12.19	8.95	8.53			
	MAMMA1002662	6.09	7.03	11.38	9.76	5.9	8.37	9.45	4.05	7.02			
5	MAMMA1002665	45.5	69.25	89.49	54.25	62.78	56.03	66.39	47.78	71.4			
	MAMMA1002671	18.2	21.13	33.24	18.58	22.22	20.8	25.58	15.85	14.12			
	MAMMA1002673	11.32	15.65	15.97	11.87	13.24	9.54	13.67	9.79	13.55			
	MAMMA1002684	12.93	20.62	16.77	15.69	20.34	11.67	14.21	4.61	5.68			
	MAMMA1002685	5.98	5.53	9.57	4.13	4.23	3.62	3.38	2.66	3.46	*	*	-
10	MAMMA1002692	8.9	8.42	9.87	7.25	7.35	7.75	2.28	5.77	1.68	*	*	-
	MAMMA1002693	9.54	10.93	23.31	11.25	9.77	9.02	7.69	7.85	6.66			
	MAMMA1002698	6.28	7.75	12.56	8.05	7.51	8.6	5.96	6.56	6.22			
	MAMMA1002699	5.33	9.37	9.52	5.1	10.31	6.85	7.2	5.75	6.86			
	MAMMA1002701	26.1	32.61	44.74	24.49	33.98	29.72	36.76	18.9	37.16			
	MAMMA1002708	10.78	13.98	19.02	10.41	15.42	12.16	15.37	13.13	12.91			
15	MAMMA1002711	14.75	23.34	34.41	24.21	29.1	25.59	25.28	14.17	20.72			
	MAMMA1002712	11.97	12.21	14.44	7.62	10.08	13.25	6.69	8.26	7.29	**	-	
	MAMMA1002716	4.68	7.56	7.57	4.77	4.14	4.42	3.66	3.97	3.59	*	-	
	MAMMA1002721	14.39	14.75	20.55	11.28	10.99	13.97	10.36	11.19	14.35			
	MAMMA1002723	7.42	10.89	16.39	7.15	8.24	4.7	4.46	4.93	6.1			
	MAMMA1002727	5.86	7.13	17.65	5.65	6.92	6.82	4.72	3.49	6.01			
20	MAMMA1002728	203.39	279.55	292.72	297.21	297.38	218.05	240.28	159.35	257.9			
	MAMMA1002742	14.02	17.19	15.98	10.59	10.28	10.18	5.71	6.31	6.17	**	**	-
	MAMMA1002743	13.89	22.29	19.83	22.11	23.47	15.54	12.39	9.78	3.07			
	MAMMA1002744	41.83	52.92	97.53	82.43	49.67	62.65	35.95	33.09	52.81			
	MAMMA1002746	1.97	3.26	4.93	1.96	3.41	2.27	2.78	1.33	2.98			
	MAMMA1002748	7.49	8.51	12.58	5.49	4.9	6.1	6.53	3.04	7			
25	MAMMA1002754	14.85	17.76	22.12	11.35	12.08	11.06	11.12	5.72	16.48	*	-	
	MAMMA1002758	3.59	1.64	11.4	2.56	2.22	2.31	1.99	1.02	1.03			
	MAMMA1002762	132.2	159.47	185.48	173.16	185.47	170.91	152.23	100.87	138.71			
	MAMMA1002764	11.39	16.4	28.38	10.76	16.28	13.47	14.81	9.65	13.58			
	MAMMA1002765	7.34	7.13	16.56	5.42	10.33	6.06	7.41	3.51	6.05			
	MAMMA1002769	10.29	13.85	24.43	5.77	11.7	10.08	4.64	4.85	10.59			
30	MAMMA1002771	3.01	3.94	5.73	4.17	3.21	3.23	5.9	1.8	2.75			
	MAMMA1002775	37.45	33.73	58.09	84.46	48.03	62.49	34.89	23.26	53.34			
	MAMMA1002780	5.07	6.15	6.67	4.47	3.43	3.09	4.23	2.8	2.68	*	*	-
	MAMMA1002782	5.1	4.96	13.58	4.26	5.86	4.79	6.17	2.83	4.75			
	MAMMA1002795	4.5	6.79	10.64	2.77	4.7	2.89	3.8	0.29	3.67			
	MAMMA1002796	5.94	10.22	11.38	3.51	7	2.99	1.75	1.74	3.7	*	-	
35	MAMMA1002805	7.95	9.3	18.26	5.36	11.22	10.66	9.08	3.61	6.4			
	MAMMA1002806	3.62	2.38	6.99	3.71	4.26	2.36	4.15	0.61	2.13			
	MAMMA1002807	18.13	22.17	39.72	27.6	21.23	23.03	24.44	17.51	32.38			
	MAMMA1002814	44.71	48.76	77.02	44.68	36.22	39.5	45.03	34.09	60.73			
	MAMMA1002817	3.99	4.65	7.4	2.69	1.42	2.37	1.9	1.38	3.31	*	-	
	MAMMA1002820	8.85	7.77	12.62	4.45	5.02	3.82	4.58	3.34	5.96	*	*	-
40	MAMMA1002830	214.03	277.57	281.23	229.96	268.12	260.12	256.75	84.59	195.17			
	MAMMA1002833	35.45	43.35	56.35	30.19	48.83	31.33	58.07	31.41	58.94			
	MAMMA1002835	2.83	4.62	14.15	2.49	4.86	1.91	2.5	0.14	2.49			
	MAMMA1002838	11.11	14.42	31.43	18.22	13.58	15.03	12.7	4.89	21.91			
	MAMMA1002842	18.2	19.62	33.31	21.65	17.53	15.65	18.5	8.81	13.46			
	MAMMA1002843	8.38	8.96	10.43	6.56	6.63	6.17	6.89	4.51	5.73	*	*	-
45	MAMMA1002844	7.03	7.49	9.64	4.13	4.73	3.32	3.78	1.87	4.09	*	*	-
	MAMMA1002845	2.68	3.31	8.62	2.22	3.45	1.79	1.92	0.92	2.1			
	MAMMA1002857	706.2	952.23	1146	1100.2	1170.5	914.39	1224.8	597.31	410.7			
	MAMMA1002858	2294.5	2940.8	1870.4	2372.6	1763.7	1167.7	3253.7	1423.1	844.59			
	MAMMA1002863	7.34	6.7	14.07	6.73	7.47	6.7	10.22	2.68	6.73			
	MAMMA1002868	37.22	42.58	112.8	66.03	44.4	45.77	34.24	23.46	47.24			
50	MAMMA1002869	27.07	27.29	44.22	74.9	47.13	42.55	24.18	21.4	25.44			
	MAMMA1002871	14.38	15.38	24.35	3.88	5.79	4.67	3.41	1.65	2.57	*	**	-
	MAMMA1002875	10.61	9.93	8.16	4.59	5.33	3.55	2.05	0.93	2.17	**	**	-
	MAMMA1002879	85.9	153.46	165.38	90.95	121.6	91.59	134.87	73.22	108.93			
	MAMMA1002880	3.99	4.47	13.13	3.43	2.59	1.91	2.37	0	2.3			
	MAMMA1002881	8.75	7.92	17.49	5.86	8.28	6.29	10.75	4.45	6.2			
55	MAMMA1002885	3.52	4.77	10.68	3.3	3.4	4.24	2.69	0.63	2.73			
	MAMMA1002886	26.73	26.24	71.16	37.73	28.68	31.72	26.26	17.17	36.38			
	MAMMA1002887	4.31	6.26	14.8	6.26	3.45	6.36	3.77	1.7	3.28			

Table 400

	MANNA1002390	16.21	20.25	27.48	13.4	13.61	14.48	13.02	8.71	18.23		
	MANNA1002392	4.77	2.83	6.02	7.9	2.39	4.12	3.41	2.08	4.6		
5	MANNA1002396	2.99	1.82	9.4	2.48	1.34	2.01	2.08	1.84	0.91		
	MANNA1002399	9.87	15.29	24.4	2.74	8.91	5.25	5.52	3.6	3.73	*	-
	MANNA1002400	72.48	90.25	98.09	48.92	68.69	49.07	35.95	25.31	45.88	* **	-
	MANNA1002409	6.49	9.18	17.47	3.97	6.62	4.36	5.84	1.59	5.43		
	MANNA1002411	9.51	7.57	20.49	9.69	12.54	10.88	11.51	6.91	7.16		
10	MANNA1002413	34.97	38.81	65.44	49.38	40.42	34.55	32.31	27.95	41.66		
	MANNA1002417	36.82	49.91	68.59	44.18	31.29	40.31	29.78	24.72	27.25		
	MANNA1002427	6.8	10.05	17.81	10.58	7.31	11.82	13	6.65	9.6		
	MANNA1002428	5.09	6.45	11.72	5.54	5.32	4.74	5.8	2.19	4.39		
	MANNA1002433	14.18	24.47	33.83	12.49	14.56	9.41	13.05	9.04	12.93		
	MANNA1002434	4.26	3.86	13.18	2.12	3.76	2.95	4.67	2.76	3.7		
15	MANNA1002446	8.39	8.67	27.55	10.82	15.66	10.5	11.46	8.07	9.95		
	MANNA1002447	4.71	6.42	13.3	4.25	7.94	6.39	4.34	1.12	6.19		
	MANNA1002454	3.91	4.62	7.05	7.12	5.04	6.67	4.66	2.3	3.21		
	MANNA1002461	27.3	34.86	56	30.05	20.73	23.5	29.37	22.7	32.72		
	MANNA1002463	9.24	10.49	15.67	10.5	6.07	6.24	8.27	4.83	7.57		
	MANNA1002464	3.98	2.85	5.66	1.75	3.24	1.37	1.77	2.41	1.96		
20	MANNA1002466	4.86	7.93	15.75	8.06	8.89	6.34	6.75	3.95	4.7		
	MANNA1002470	4.68	6.3	14.57	5.95	13.1	4.89	3.64	2.83	3.84		
	MANNA1002475	17.39	22.9	40.72	24.51	29.14	16.55	20.39	12.61	16		
	MANNA1002480	51.48	94.76	100.05	63.97	58.87	37.35	60.56	13.3	33.46		
	MANNA1002485	19.11	15.89	43.85	26.44	20.15	19.25	16.32	13.38	18.32		
	MANNA1002494	22.86	28.77	50.5	24.3	23.02	18.38	23.12	24.83	27.28		
25	MANNA1002498	5.27	2.99	4.95	1.99	2.08	1.95	2.18	0.89	1.64	*	-
	MANNA1002524	11.96	11.09	24.02	7.72	7.78	5.46	10.41	6.02	6.07	*	-
	MANNA1002530	277.32	388.85	679.97	473.31	550.87	394.12	467.12	251.99	328.81		
	MANNA1002538	1.65	4.73	10.72	1.52	2.62	2.66	2.05	0.61	2.49		
	MANNA1002545	2.59	2.37	13.88	1.77	4.03	1.39	1.68	0.09	1.93		
30	MANNA1002554	115.83	185.59	273.08	258.19	207.86	207.1	180.88	93.96	83.88		
	MANNA1002556	4.03	6.04	7.54	5.06	3.68	5.28	2.7	3.76	3.11		
	MANNA1002561	15.67	18.37	24.06	5.72	9.69	8.96	6.7	7.18	2.52	*	-
	MANNA1002565	20.13	19.37	64.78	24.22	20.52	19.19	20.48	14.5	21.19		
	MANNA1002566	5.49	3.26	6.28	2.19	3.2	3.17	4.16	1.97	2.42		
	MANNA1002571	12.97	18.44	36.1	16.95	12.11	12.28	16.66	10	13.05		
	MANNA1002573	4.02	5.04	11.33	1.38	3.48	2.19	3.22	1.15	2.21		
35	MANNA1002576	13.78	23.79	37.39	16.06	19.72	15.26	13.9	8.77	10.92		
	MANNA1002584	9.21	12.39	27.02	16.71	13.29	9.15	13.96	6.48	12.24		
	MANNA1002585	6.36	5.36	13.56	11.07	9.12	8.7	7.18	4.4	6.71		
	MANNA1002586	21.75	21.37	47.64	26.29	19.9	20.88	19.88	14.32	18.17		
	MANNA1002589	31.16	39.3	81.02	35.29	36.93	32.17	47.24	33.3	29.17		
	MANNA1002590	8.65	6.16	15.55	9.32	8.84	9.26	5.06	3.24	4.01		
40	MANNA1002593	8.47	7.53	30	8.1	8.37	7.45	8.63	8.66	8.47		
	MANNA1002597	11.34	16.94	30.07	13.1	12.76	11.04	12.91	7.38	12.5		
	MANNA1002598	9.23	11.39	26.12	15.11	13.9	10.6	14.04	4.47	10.54		
	MANNA1002603	7.01	8.63	19.81	9.42	12.79	9.69	13.58	3.61	8.05		
	MANNA1002612	0.64	0.54	1.4	4.76	2.04	2.86	2.1	0	0.92		
	MANNA1002617	1.99	2.39	4.19	2.19	2.2	2.69	2.8	1.84	1.34		
45	MANNA1002618	16.06	18.63	46.52	25.87	25.59	22.23	19.52	10.7	17.56		
	MANNA1002619	7.62	6.62	18.84	13.61	11.83	9.5	6.72	6.93	5.26		
	MANNA1002622	3.06	2.72	4.49	4.32	2.96	14.63	2.48	0.95	1.91		
	MANNA1002623	2.38	2.14	3.99	2.64	2.39	1.68	1.62	0	1.12		
	MANNA1002625	158.57	183.87	332.03	71.22	106.04	287.72	518.89	291.59	178.62		
	MANNA1002627	6.48	11.48	14.86	11.21	19	9.56	12.93	6.03	9.03		
50	MANNA1002629	20.19	19.33	33.94	25.61	18.48	21.46	13.72	14.18	19.09		
	MANNA1002631	4.6	4.73	9.76	3.98	2.11	3.82	1.12	0.35	1.47	*	-
	MANNA1002633	64.25	69.26	104.49	122.15	95.76	107.58	86.41	60.3	67.07		
	MANNA1002636	30.48	34.84	49.63	34.43	30.9	28.96	42.01	28.77	46.58		
	MANNA1002637	2.95	5.14	7.06	4.15	3.68	2.57	4.18	1.82	4.8		
	MANNA1002646	3.86	4.18	6.8	3.4	4.56	2.51	2.73	1.37	2.49		
55	MANNA1002648	136.47	120.19	147.15	150.69	135.33	106.16	124.23	86.88	146.01		
	MANNA1002650	1.26	1.85	1.16	1.57	0.72	1.64	0.26	0.46	0.54	*	-
	MANNA1002652	32.12	44.28	46.88	48.41	32.58	36.9	24.05	25.91	28.22	*	-

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	MAMMA1002145	59.23	60.84	77.53	26.62	46.01	14.01	21.49	27.44	27.68	*	**	-	-
	MAMMA1002147	5.58	4.1	8.16	3.48	3.17	3.48	3.61	1.58	1.54				
5	MAMMA1002153	2.82	4.71	7.06	3.26	5.54	4.01	3.33	4.14	1.01				
	MAMMA1002155	2.48	3.25	10.86	2.89	3.23	2.81	4.53	1.33	2.05				
	MAMMA1002156	1.89	5.51	11.57	0.4	3.84	3.05	2.17	1.45	3.44				
	MAMMA1002158	31.98	68.42	55.01	8.28	17.41	13.99	16.74	7.42	4.4	*	*	-	-
	MAMMA1002164	1.74	3.48	10.17	3.33	5.39	3.9	5.07	0.22	3.21				
10	MAMMA1002165	14.96	21.57	31.25	20.06	14.73	17.78	14.07	10.69	17.29				
	MAMMA1002170	22.65	29.54	35.22	10.64	9.2	7.67	9.57	17.6	7.86	**	*	-	-
	MAMMA1002174	5.6	8.95	10.45	5.64	4.66	7.45	4.49	3.82	4.25		*	-	-
	MAMMA1002175	51.5	30.74	138.92	91.4	93.51	53.96	49.64	44.62	29.88				
	MAMMA1002180	26.41	35.57	64.21	33.73	28.48	23.51	38.7	27.33	38.33				
	MAMMA1002198	37.99	80.63	138.79	18.29	47.55	46.05	13.24	10.75	8.83				
15	MAMMA1002205	15.14	20.56	52.79	21.27	16.14	13.46	24.47	10.14	16.64				
	MAMMA1002206	2.93	3.59	10.31	2.52	3.96	2.18	3.53	0.98	3.24				
	MAMMA1002209	183.5	268.42	348.13	282.08	352.1	269.26	189.69	127.24	124.82				
	MAMMA1002215	24.38	31.33	40.05	29.39	28.67	24.47	25.62	23.02	20.25				
	MAMMA1002219	2.71	0.94	3.73	1.84	2.07	2.08	0.86	0.6	1.35				
	MAMMA1002224	15.23	8.24	19.36	12.34	12.12	11.67	6.61	10.38	6.31				
20	MAMMA1002229	16.46	14.44	27.79	9.51	10.83	11.84	7.32	11.92	8.59				
	MAMMA1002230	22	37.1	57.12	14.15	20.28	20.13	22.25	15.46	15.8				
	MAMMA1002233	70.38	89.47	122.68	8.19	33.96	39.69	218.52	154.84	147.67	*	*	-	+
	MAMMA1002234	7.92	8.85	12.79	4.92	10.39	5.68	11.44	4.89	8.58				
	MAMMA1002236	136.07	148.16	220.88	36.8	57.62	87.11	34.48	55.67	55.33	*	*	-	-
	MAMMA1002243	20.02	20.33	38.87	38.33	25.12	25.3	21.56	17.63	24.13				
25	MAMMA1002250	2.68	3.81	6.67	5.79	2.6	4.11	4.54	2.26	2.04				
	MAMMA1002253	0.57	1.7	6.1	2.21	1.65	3.98	2.62	1.41	1.86				
	MAMMA1002267	9.29	9.06	17.68	10.28	11.75	12.03	8.49	4.34	11.22				
	MAMMA1002268	0.57	3.72	6.93	16.18	7.45	4.5	3.59	1.73	3.92				
	MAMMA1002269	1.7	2.24	4.35	2.97	8.98	8.51	2.83	1.4	3.35				
	MAMMA1002282	2.86	3.76	4.32	2.98	7.03	4.77	3.56	5.53	5.39				
30	MAMMA1002292	15.43	25.26	45.69	33.39	19.64	22.84	15.48	15.23	21.7				
	MAMMA1002293	3.2	3.94	4.4	4.2	3.49	3.24	3.79	4.35	3.43				
	MAMMA1002294	9.41	6.68	12.5	3.33	7.99	10.5	5.31	8.44	4.21				
	MAMMA1002297	5.35	6.59	11.35	7.95	7.71	8	2.76	2.6	3.09				
	MAMMA1002298	3.44	2.01	8.19	4.11	4.61	4.39	2.48	2.96	3.08				
	MAMMA1002299	2.99	4.55	14.53	7.08	6.73	6.49	4.69	0.99	4.04				
35	MAMMA1002308	5.47	9.33	18.02	10.01	8.02	7.61	7.23	3.03	7.9				
	MAMMA1002310	9.13	10.94	18.01	13.86	17.88	11.7	14.2	7.14	10.5				
	MAMMA1002311	10.67	15.07	18.09	4.83	5.07	6.09	1.95	3.51	2.65	*	**	-	-
	MAMMA1002312	30.9	42.06	49.45	34.33	28.66	29.57	22.98	30.71	35.58				
	MAMMA1002317	5.72	5.28	10.29	3.9	3.7	4.94	4.31	4.41	4.7				
	MAMMA1002319	11.12	14.39	20.39	9.58	9.59	6.6	12.89	13.6	13.85				
40	MAMMA1002322	2.09	3.24	7.26	1.71	2.86	1.28	1.75	1.09	1.05				
	MAMMA1002329	17.33	19.74	32.38	18.64	18.47	17.53	19	10.56	17.8				
	MAMMA1002332	2.58	4.23	6.84	4.89	4.12	2.94	2.37	0.43	2.61				
	MAMMA1002333	1.45	2.08	7.07	4.55	5.97	6.64	2.39	0.64	2.14				
	MAMMA1002335	12.65	18.48	23.72	7.59	6.81	10.8	3.17	6.12	8.64	*	*	-	-
	MAMMA1002339	7.58	16	19.05	10.2	7.64	10.35	7.33	10.1	11.23				
45	MAMMA1002347	10.15	21.39	24.74	8.5	9.44	11.14	10.35	10.5	16.39				
	MAMMA1002351	7.61	10.35	21.39	9.12	8.67	7.13	7.44	4.46	10.65				
	MAMMA1002352	3.96	6.91	9.29	4.01	3.84	4.53	2.75	1.39	5.61				
	MAMMA1002353	8.39	10.78	23.6	6.39	12.46	12.35	7.8	9.12	7.88				
	MAMMA1002355	25.93	29.71	53.72	28.58	30.76	23.64	29.82	21.55	27.08				
	MAMMA1002356	28.68	42.92	56.16	40.53	46.92	35.69	34.66	27.13	37.63				
50	MAMMA1002359	9.7	13.9	19.55	15.93	13.24	12.52	6.92	6.02	14.49				
	MAMMA1002360	85.31	125.75	125.12	50.2	80.78	78.44	18.48	22.03	30.83	**		-	
	MAMMA1002361	4.68	5.33	13.4	6.92	5.11	6.08	3.35	2.61	7.85				
	MAMMA1002362	4.53	5.8	12.93	6.01	7	5.37	5.23	3.68	7.06				
	MAMMA1002367	13.67	17.12	24.55	7.42	11.11	9.42	8.07	3.62	6.96	*		-	
	MAMMA1002371	2.82	3.35	9.8	0.89	2.2	2.61	2.45	2.44	1.99				
55	MAMMA1002380	7.87	13.26	17.64	9.95	9.26	6.58	10	7.71	10.46				
	MAMMA1002384	91.44	168.34	87.56	61.83	126.23	76.47	56.65	15.61	10.22	*		-	
	MAMMA1002385	2.04	1.77	7.54	5.82	2.53	3.4	1.44	1.54	1.62				

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	MAMMA1001851	3.04	3.08	4.42	3.11	3.54	2.54	2.52	0.56	2.38		
	MAMMA1001852	4.14	3.26	11.19	3.61	3.33	5.4	1.88	2.12	2.02		
5	MAMMA1001854	1.51	2.81	6.9	2.87	1.88	2.16	2.29	1.4	2.53		
	MAMMA1001858	19.51	21.77	35.8	19.05	16.81	16.88	15.59	11.41	20.06		
	MAMMA1001864	13.26	17.9	34.07	19.43	19.24	16.98	19.91	10.97	23.39		
	MAMMA1001868	36.41	51.95	37.39	27.02	32.06	24.02	38.47	28.02	40.25		
	MAMMA1001874	10.43	12.98	12.56	15.09	16.33	15.06	13.47	8	7.34	*	+
10	MAMMA1001878	8.74	13.09	17.2	10.72	7.64	9.1	6.72	5.47	7.12		
	MAMMA1001880	4.07	4.26	5.15	5.16	2.51	4.87	2.76	4.1	3.13		
	MAMMA1001885	11.53	16.59	20.89	11.02	9.99	7.12	10.84	10.3	12.47		
	MAMMA1001890	18	26.55	27.35	18.41	24.49	18.88	18.85	18.1	22.42		
	MAMMA1001893	9.74	13.29	21.33	8.06	11.26	12.34	13.42	8.15	13.42		
	MAMMA1001901	5.91	10.54	9.81	6.07	8.49	8.27	5.4	3.05	2.51	*	-
15	MAMMA1001907	2.61	3.62	7.04	35.62	7.01	7.16	2.77	1.6	2.87		
	MAMMA1001908	8.52	7.37	12.33	6.04	6.79	5.83	5.18	1.69	2.73	*	-
	MAMMA1001919	28.87	31.67	48.08	80.53	41.26	66.58	17.81	21.06	28		
	MAMMA1001931	5.05	3.95	9.8	4.52	3.24	2.79	3.04	5.51	2.52		
	MAMMA1001937	14.26	19.97	17.05	11.88	9.67	16.46	11.3	11.49	14.97		
	MAMMA1001951	10.03	11.71	22.7	11.88	9.84	10.7	11.47	12.01	12.62		
20	MAMMA1001956	993.12	926.97	1643.3	1427.1	1487.9	1519.7	1062.9	902.24	975.35		
	MAMMA1001957	5.28	8.16	19.85	8.48	7.2	6.6	6.19	4.02	7.97		
	MAMMA1001960	4.45	5.22	11.99	3.94	5.66	4.15	4.01	2.89	2.74		
	MAMMA1001963	9.87	14.96	20.88	14.68	14.73	14.4	8.07	6	3.43		
	MAMMA1001969	11.23	14.81	19.75	14.09	10.96	16.21	10.31	8.8	19.07		
	MAMMA1001970	4.02	5.46	5.52	3.48	4.15	19.31	4.95	4.07	2.18		
25	MAMMA1001978	13.78	19.86	26.28	16.09	13.28	16.86	13.67	12.24	23.16		
	MAMMA1001992	6	4.83	12.16	2.11	5.12	11.04	3.79	2.55	5.99		
	MAMMA1001994	4.27	7.76	10.55	16.33	8.69	5.41	3.19	4.78	5.52		
	MAMMA1002008	2.62	3.39	13.27	3.37	3.96	2.92	2.97	2.86	1.68		
	MAMMA1002009	3.5	5.4	13.89	4.02	6.32	3.6	2.53	1.71	1.09		
	MAMMA1002011	3.21	3.57	5.26	2.05	5.5	3.92	2.9	2.56	2.89		
30	MAMMA1002022	23.41	32.84	43.13	18.55	24.01	20.69	18.94	17.16	21.03		
	MAMMA1002024	4.69	8.48	7.51	4.82	3.7	4.18	3.88	2.33	4.09		
	MAMMA1002032	22.72	24.3	34.55	17.1	16.95	18.46	20.89	15.73	25.86		
	MAMMA1002033	5.27	4.69	9.29	4.37	4.11	4.75	8.94	7.94	5.81		
	MAMMA1002041	6.92	11.13	16.18	4.7	5.73	4.73	2.24	3.33	5.35		
	MAMMA1002042	4.21	5.9	12.97	1.93	6.7	3.65	5.79	7.29	5.48		
35	MAMMA1002045	3.58	4.86	12.3	4.27	5.48	4.41	3.9	4.78	2.67		
	MAMMA1002047	6.8	7.34	9.55	7.31	12.68	11.46	8.75	3.44	4.92		
	MAMMA1002056	2.57	2.86	6.49	3.74	3.21	3.7	1.38	1.89	2.78		
	MAMMA1002058	12.8	12.32	19.01	10.51	10.24	9.8	10.2	7.07	10.6		
	MAMMA1002060	3.44	2.98	10.21	3.91	3.36	3.73	4.39	1.38	3.75		
	MAMMA1002065	5.77	6.94	12.55	3.81	3.34	3.54	3.78	2.53	2.66		
40	MAMMA1002068	5.43	5.26	11.94	2.41	4.51	4.09	4.21	2.57	4.68		
	MAMMA1002070	3.38	5.43	10.65	2.42	2.8	1.57	1.88	0.34	3.17		
	MAMMA1002078	3.03	3.94	12.3	1.63	3.98	2.25	2.32	1.3	2.65		
	MAMMA1002080	11.92	16.79	18.92	3	9.03	5.76	7.02	2.85	4.16	*	**
	MAMMA1002082	8.32	5.53	12.97	8.62	3.65	5.55	2.81	1.42	4.34		
	MAMMA1002084	5.95	5.89	9.82	6.01	4.3	5.5	3.1	3.76	3.18	*	-
45	MAMMA1002087	4.18	2.51	5.36	3.1	3.55	1.79	2.57	2.24	1.51		
	MAMMA1002091	4.77	6.45	9.77	6.57	5.69	2.84	2.82	2.99	2.11	*	-
	MAMMA1002093	7.77	50.44	12.78	4.52	8.15	3.8	10.66	4.49	6.61		
	MAMMA1002095	2.07	3	10.99	1.29	1.85	0.7	1.89	0.49	1.88		
	MAMMA1002108	11.08	15.62	32.13	20.98	14.75	19.73	14.58	13.23	20.21		
50	MAMMA1002112	11.23	16.95	26.2	16.44	15.29	9.14	16.6	9.46	11.52		
	MAMMA1002118	15.81	20.81	31.7	25.23	19.06	12.25	11.43	15.5	24.54		
	MAMMA1002119	10	9.39	20.1	14.99	8.54	8.01	7.06	7.05	10.3		
	MAMMA1002125	8.81	6.62	15.53	7.95	7.53	5	5.63	5.81	8.41		
	MAMMA1002126	38.45	22.62	47.44	8.75	23.38	14.29	26.49	8	13.65		
	MAMMA1002128	9.97	18.56	26.22	10.58	12.77	9.47	19.63	7.42	11.63		
	MAMMA1002132	6.21	7.25	34.7	5.61	8.67	4.88	6.36	3.29	9.25		
55	MAMMA1002140	16.11	29.14	22.82	11.98	12.1	5.91	9.18	3.59	4.66	*	*
	MAMMA1002142	3.45	5.23	11.28	4.42	6.67	3.75	4.53	0.93	3.45		
	MAMMA1002143	16.35	16.97	31.08	20.92	14.17	21.18	9.45	9.06	16.49		

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	MANMA1001606	37.25	38.33	44.14	13.41	26.09	29.22	7.5	16	15.16	*	**	-	-
5	MANMA1001609	2.18	4.53	5.97	1.87	1.39	1.8	4.17	2.27	1.51				
	MANMA1001614	10.91	17.85	21.09	12.42	12.51	13.02	16.85	9.92	19.29				
	MANMA1001615	1.95	3.17	11.55	2.77	2.24	1.35	2.33	1.46	2.06				
	MANMA1001619	11.51	16.15	15.55	6.34	13.88	8.59	8.19	7.08	5.14	*		-	
	MANMA1001620	4.59	6.42	11.46	1.77	3.88	3.43	3.11	1.71	4				
	MANMA1001623	46.36	31	53.81	63.16	42.72	47.15	31.15	25.06	25.66				
10	MANMA1001626	23.59	24.5	38.75	33.26	30.12	18.9	7.93	11.89	22.41				
	MANMA1001627	23.89	43.37	29.47	17.73	21.82	12.9	6.43	10	7.98	*		-	
	MANMA1001630	9.27	13.78	11.56	7.25	5.63	6.82	2.81	4.06	2.66	*	**	-	-
	MANMA1001633	3.1	2.16	7.03	2.23	2.28	2.16	0.8	0.79	2.49				
	MANMA1001634	3.53	6.28	9.01	3.76	4.19	2.88	3.9	2.94	2.79				
	MANMA1001635	3.28	4.64	15.97	2.28	2.56	1.53	6.22	1.65	2.6				
15	MANMA1001649	3.05	5.72	15.55	3.26	4.9	2.56	4.26	1.39	3.94				
	MANMA1001654	4.23	5.9	13.55	3.26	5.84	3.8	3.63	1.4	2.71				
	MANMA1001660	39.09	42.24	88.43	56.56	52.47	44.31	41.47	39	60.86				
	MANMA1001663	8.52	11.06	11.71	10.2	6.57	5.19	7.96	6.94	8.33				
	MANMA1001670	5.91	7.75	6.72	4.07	2.57	5.22	5.32	3.26	4.27	*	*	-	-
	MANMA1001671	56.61	62.54	66.07	56.8	37.97	53.72	50.14	43.96	58.96				
20	MANMA1001679	6.69	13.86	10.25	4.05	5.47	3.58	4.46	7.43	6.03				
	MANMA1001683	15.6	21.19	31.72	12.85	17.62	10.3	18.14	10.49	14.57				
	MANMA1001686	4.11	3.45	11.77	4.66	4.36	4.4	4.89	1.58	2.59				
	MANMA1001688	4.96	5.41	15.23	3.79	4.7	4.42	2.65	1.46	3.58				
	MANMA1001689	28.44	25.89	32.18	7.85	17.39	15.22	8.44	8.89	20.79	*	*	-	-
	MANMA1001692	5.97	3.79	10.24	3.42	9.67	3.1	3.28	4.26	5.93				
25	MANMA1001711	4.75	4.66	8.45	3.21	3.28	2.22	2.67	8.04	0.88				
	MANMA1001715	4.22	4.73	9.43	3.9	2.04	3.47	3.76	4.11	2.61				
	MANMA1001730	3.39	2.22	8.02	1.72	1.43	1.79	1.71	1.46	1.86				
	MANMA1001735	8.81	9.96	22.03	12.16	11.12	10.37	6.84	5.43	7.1				
	MANMA1001740	2.28	4.25	11.56	2.64	4.24	1.86	2.43	1.41	3.14				
	MANMA1001743	10.52	18.5	19.92	5.55	7.23	6.37	6.45	1.87	3.83	*	*	-	-
30	MANMA1001744	4.59	3.07	5.75	3.86	4	4.16	2.05	2.19	3.63				
	MANMA1001745	12.62	11.29	21.77	12.13	11.56	10.5	13.92	9.55	13.74				
	MANMA1001751	11.11	8.28	15.68	3.47	4.56	3.38	2.93	2.2	1.66	*	*	-	-
	MANMA1001752	5.63	3.02	9.77	2.83	5.25	2.56	2.97	2.42	0.97				
	MANMA1001754	23.39	32.69	51.88	31.96	35.9	26.06	34.24	24.41	25.71				
	MANMA1001757	5.44	8.41	15.4	4.99	7.7	6.28	7.54	3.86	5.12				
35	MANMA1001760	2.58	5.03	10.84	4.07	3.09	11.35	3.87	1.47	1.67				
	MANMA1001764	1.11	5.18	12.27	3.88	5.02	2.55	4.09	0.44	2.15				
	MANMA1001767	3.1	2.25	5.04	3.83	2.36	7.31	2.64	1.02	1.81				
	MANMA1001768	11.92	10.35	26.44	10.9	12.51	10.66	12.91	3.89	9.69				
	MANMA1001769	14.67	15.11	32.18	14.99	14.79	13.17	17.74	10.66	13.05				
	MANMA1001771	5.81	5.01	9.7	5.52	6.04	4.42	3.1	4.15	2.2				
40	MANMA1001773	40.73	46.88	88.4	41.31	38.76	36.3	25.57	21.1	22.63				
	MANMA1001778	15.44	10.48	54.31	14.07	14.31	18.11	19.19	12.85	7.66				
	MANMA1001783	12.5	12.45	33.56	11.62	10.75	17.89	7.63	5.83	3.14				
	MANMA1001785	4.08	3.99	8.35	2.35	3.38	3.07	3.75	1.41	2.66				
	MANMA1001788	6.1	6.55	20.74	18.01	10.68	15.67	7.37	4.2	7.06				
	MANMA1001790	1.31	0.72	1.87	1.36	1.35	1.32	1.67	0.59	1.36				
45	MANMA1001800	2.19	1.86	2.06	3.44	3.43	3.44	1.48	0.44	1.31	**	*	+	-
	MANMA1001804	9.74	13.74	26.09	23.09	17.83	16.06	15.63	9.04	8.32				
	MANMA1001806	15.42	15.2	24.16	21.08	18.33	17.7	13.08	11.61	12.4				
	MANMA1001812	2.66	2.87	5.53	5.25	2.56	2.24	1.76	1.16	2				
	MANMA1001815	0.82	1.33	4.14	2.35	1.64	0.9	2.33	0	1.04				
	MANMA1001817	7.15	8.39	26.7	8.94	11.15	9.35	10.6	5.97	8.95				
50	MANMA1001818	11.74	18.24	23.68	11.92	10.04	13.23	8.57	10.06	9.02				
	MANMA1001819	15.72	24.34	43.45	17.35	11.3	16.16	12.64	12.63	16.41				
	MANMA1001820	22.82	19.64	37.64	42.37	24.16	31.98	15.13	16.66	19.01				
	MANMA1001824	8.99	12.77	18.28	12.69	14.09	15.06	12.72	9.76	11.95				
	MANMA1001832	7.22	10.73	12.96	9.65	7.55	9.34	6.24	4.93	7.88				
	MANMA1001836	9.42	23.13	19.47	21.69	15.29	10.85	11.8	5.71	11.11				
55	MANMA1001837	3.6	4.29	8.23	5.89	6.7	8.36	4.62	1.11	4.3				
	MANMA1001848	16.38	27.46	23.83	25.7	38.67	49.58	24.68	13.45	27.7				
	MANMA1001850	4.58	4.63	5.94	6.14	2.91	3.96	4.34	2.05	2				

Table 396

5	MAMMA1001286	29.08	30.31	35.52	8.75	7.01	14.48	10.18	16.99	22.31	**	*	-	-
	MAMMA1001289	8.87	7.96	10.4	9.73	7.53	9	7.31	3.62	7.73				
	MAMMA1001292	24.97	44.07	66.54	9.34	36.59	35.11	13.77	28.43	10.4				
	MAMMA1001296	4	4.17	8.01	2.8	3.49	2.38	2.75	2.82	4.19				
	MAMMA1001298	7.53	9.89	10.46	2.88	6.46	2.62	5.34	3.84	1.86	*	*	-	-
	MAMMA1001305	3.74	4.35	13.81	2.61	4.06	3.12	3.76	5.53	2.5				
	MAMMA1001309	10.42	15.14	31.52	12.87	13.99	9.19	10.23	3.68	9.96				
10	MAMMA1001310	2.92	5.46	13.25	3.59	3.89	1.96	3.03	0.43	1.21				
	MAMMA1001322	2.17	2.64	6.93	3.3	2.91	2.5	1.12	1.37	2.03				
	MAMMA1001324	4.53	3.59	5.55	3.63	2.78	3	2.65	1.6	1.68	*		-	
	MAMMA1001330	4.9	5.1	12.68	7.84	4.19	4.82	3.36	2.64	3.18				
	MAMMA1001333	4.95	3.16	7.58	2.56	3.62	3.33	2.38	2.25	1.89				
	MAMMA1001334	31.92	25.66	43.45	18.83	15.13	18.08	17.92	22.66	13.99	*		-	
15	MAMMA1001337	4.26	3.84	13.74	3.62	4.64	5.34	4	3.29	3.16				
	MAMMA1001341	4.05	4.96	11.41	4.38	5.21	4.83	5.24	4.52	3.64				
	MAMMA1001343	2.73	5.65	10.8	3.22	4.68	4.2	3.94	1.64	3.95				
	MAMMA1001344	2.25	2.47	7.73	4.27	4.36	3.62	2.73	1.3	3.12				
	MAMMA1001346	5.84	6.71	17.37	10.65	9.35	8.33	6.74	3.26	6.06				
20	MAMMA1001383	6.99	8.47	18.39	14.27	8.27	9.89	7.45	4.12	6.12				
	MAMMA1001388	25.28	28.84	63.23	32.5	23.98	25.52	24.7	25.15	17.69				
	MAMMA1001396	15.94	20.37	30.49	14.54	24.62	26.46	14.45	46.08	11.25				
	MAMMA1001397	14.75	18.37	41.81	21.11	18.82	16.79	17.96	11.5	21.37				
	MAMMA1001401	2.94	3.12	6.56	1.21	2.62	3.02	2.46	0.76	2.89				
	MAMMA1001408	1.5	3.57	7.4	2.81	1.88	3.96	5.62	0.76	2.46				
	MAMMA1001411	6.08	3.93	9.81	9.46	6.06	8.29	3.36	4.18	4.81				
25	MAMMA1001414	6.51	7.32	14.7	23.79	6.86	10.67	5.51	4.82	6.45				
	MAMMA1001415	3.95	8.97	8.88	5.52	5.96	8.18	6.06	2.79	4.69				
	MAMMA1001418	8.77	9.1	12.47	12.83	10.06	11.32	13.03	7.4	7.63				
	MAMMA1001419	7.73	7.93	11.48	9.88	6.73	10.35	9.18	5.19	6.18				
	MAMMA1001420	5.14	4.76	6.97	10.23	6.22	6.56	5.34	3.95	4.64				
	MAMMA1001426	5.19	7.71	9.63	5.55	8.11	6.73	7.6	4.91	8.87				
30	MAMMA1001428	3.64	3.18	7.92	6.62	3.95	5.99	3.35	1.43	3.18				
	MAMMA1001432	12.35	13.88	21.83	18.83	11.53	18.33	8.1	11.28	14.53				
	MAMMA1001435	20.05	30.02	44.24	21.04	23.86	21.39	14.97	14.52	16.51				
	MAMMA1001442	6.94	7.75	11.08	5.53	5.57	7.7	6.68	4.54	7.44				
	MAMMA1001446	26.48	31.11	44.32	20.63	23.27	24.39	8.85	5.14	8.85	**		-	
	MAMMA1001450	11.55	15.7	15.65	12.27	11.07	10.5	14.48	8.46	14.95				
35	MAMMA1001452	4.86	7.91	8.36	9.2	4.08	6.24	7.72	3.93	6.05				
	MAMMA1001465	8.92	12.52	15.39	11.83	13.04	10.87	16.92	7.42	13.06				
	MAMMA1001476	132.51	144.85	285.99	418.67	194.96	350.68	82.5	94.08	100.24				
	MAMMA1001478	18.64	21.54	45.82	21.72	20.69	19.34	14.71	16.9	28				
	MAMMA1001479	11.41	20.27	19.2	13.74	10.72	9.96	13.4	11.81	15.33				
	MAMMA1001487	48.7	29.91	87.89	41.22	44.32	54.61	29.8	35.15	13.96				
40	MAMMA1001498	4.35	2.96	9.82	3.07	3.33	1.99	1.92	4.02	3.09				
	MAMMA1001501	4	5.27	10.62	3.92	4.81	5.08	4.06	2.78	3.9				
	MAMMA1001502	3.5	3.68	8.07	3.61	4.35	4.14	3.43	1.95	7.01				
	MAMMA1001510	11.46	20.55	16.62	13.12	12.05	13.04	13.33	7.96	10.37				
	MAMMA1001522	4.88	10.87	13.99	8.11	4.33	5	2.46	3.78	5.04				
	MAMMA1001529	59.97	69.9	94.57	30.63	46.88	57.69	31.84	37.9	51.83	*		-	
45	MAMMA1001532	28.38	37.51	51.46	23.78	26.52	23.49	26.6	24.11	41.98				
	MAMMA1001533	19.86	8.23	26.5	6.01	12.03	8.61	7.93	4.83	6.28				
	MAMMA1001534	11.45	13.72	20.66	6.58	8.79	8.61	9.6	5.71	8.05				
	MAMMA1001535	10.07	15.11	17.7	7.42	8.38	7	2.75	2.21	3.48	*	**	-	-
	MAMMA1001547	5.79	10.79	16.77	6.18	7.64	6.73	7.59	4.04	7.84				
	MAMMA1001551	1.79	3.35	10.09	2.36	3.75	2.54	3.47	0.46	2.39				
50	MAMMA1001569	5.44	7.64	12.53	5.97	3.18	5.89	2.71	3.74	6.78				
	MAMMA1001575	19.87	31.06	32.95	20.06	20.37	18.83	18.83	16.95	22.11				
	MAMMA1001576	4.97	13.68	9.34	4.55	6.26	3.49	1.76	5.49	4.16				
	MAMMA1001584	17.12	20.22	32.05	43.71	22.66	30.09	21.3	15.63	24.89				
	MAMMA1001586	3.44	2.63	9.42	2.75	2.52	1.44	2.63	0.98	2.02				
	MAMMA1001590	4.74	3.32	9.43	0.65	2.72	2.98	5.93	1.26	2.67				
55	MAMMA1001599	6.91	10.35	19.33	5.33	6.4	5.4	7.64	5.96	7.44				
	MAMMA1001600	12.56	24.27	25.25	10.57	15.9	13.61	21.16	12.02	17.72				
	MAMMA1001604	20.93	20.87	41.04	17.28	15.57	5.46	4.43	10.16	15.17				

Table 395

	MAHMA1001030	7.4	11.72	12.89	11.66	7.79	10.11	7.28	6.15	7.3			
5	MAHMA1001035	39.22	53.03	74.6	50.1	27.34	38.76	23.51	31.74	38.44			
	MAHMA1001036	3.31	5.83	5.4	5.92	5.35	4.3	4.37	2.81	3.74			
	MAHMA1001037	24.23	31.03	53.22	27.23	25.26	31.34	28.91	17.23	25.45			
	MAHMA1001038	12.98	20.58	21.05	13.13	14.71	11.55	14.52	10.1	18.77			
	MAHMA1001041	1.54	2.15	5.75	4.24	4.69	7.8	3.02	1.22	2.08			
	MAHMA1001043	1.39	2.46	4.67	3.06	4.27	3.32	4.35	1.52	2.3			
10	MAHMA1001050	1.33	4.39	4.83	3.8	5.59	5.14	4.27	1.05	2.57			
	MAHMA1001054	19.75	24.03	50.38	32.02	21.79	27.11	13.83	16.21	20.46			
	MAHMA1001059	3.19	4.55	5.53	3.65	3.25	5.05	2.54	10.99	3.33			
	MAHMA1001066	5.02	4.16	8.84	5.66	2.95	3.72	1.82	2.25	3.08			
	MAHMA1001067	17.58	21.79	31.08	14.83	16.92	18.84	15.29	17.34	18.26			
	MAHMA1001072	2.53	1.88	3.3	1.93	1.68	2.41	2.3	1.25	2			
15	MAHMA1001073	0.84	2.27	6.85	1.01	1.41	5.24	0.75	4.07	1.61			
	MAHMA1001074	24.28	39.17	47.61	31.96	28.12	24.54	31.91	23.64	32.81			
	MAHMA1001075	1.48	5.48	8.44	3.99	4.91	2.99	2.44	2.09	3.72			
	MAHMA1001078	21.72	26.12	44.96	33.39	16.5	21.38	18.07	17.35	25.34			
	MAHMA1001080	12.04	17.72	15.6	14.26	12.06	16.7	6.39	6.97	11.08	*	-	
	MAHMA1001082	4.59	9.29	12.12	6.84	2.79	4.23	4.41	1.12	3.78			
20	MAHMA1001091	16.73	23.95	30.14	19.81	16.46	17.71	21.25	12.1	23.6			
	MAHMA1001092	7.81	11.13	15.6	7.44	8.17	6.28	7.18	4.54	7.68			
	MAHMA1001094	11.74	16.72	20.33	14.77	10.66	23.71	14.53	12.3	17.29			
	MAHMA1001105	45.26	57.55	66.73	41.25	47.01	32.75	30.09	24.22	28.58	*	-	
	MAHMA1001110	7.59	14.4	12.56	8.44	9.45	9.37	9.82	4.24	0.9			
	MAHMA1001126	37.56	44.85	64.49	49.13	34.88	37.49	24.63	36.31	49.73	*	-	
25	MAHMA1001133	5.76	9.31	8.87	3.28	3.01	4.08	2.64	5.37	6.65	*	-	
	MAHMA1001139	7.27	14.13	10.95	4.54	4.49	4.1	6.16	2.44	5.84	*	-	
	MAHMA1001141	6.69	8.3	15.1	2.9	4.24	4.06	2.47	2.3	5.32			
	MAHMA1001143	3.83	3.21	9.91	1.84	2.97	1.81	3.49	1.28	2.11			
	MAHMA1001145	2.22	12.54	14.95	2.6	3.58	6.73	2.4	2.76	3.58			
	MAHMA1001150	10.97	15.54	29.9	12.2	13.68	9.93	10.31	8.5	12.03			
30	MAHMA1001154	5.02	9.06	11.37	4.94	8.36	4.03	3.8	2.43	3.42			
	MAHMA1001159	6.91	7.68	15.13	5.63	3.35	4.72	2.34	2.78	4.72			
	MAHMA1001161	109.99	168.02	168.22	189.81	141.37	144.88	81.63	94.69	118.15			
	MAHMA1001162	7.19	5.58	7.65	7.64	5.6	5.96	4.02	2.29	5.78			
	MAHMA1001181	2.37	3.1	6.4	4.16	1.93	2.21	2.1	2.11	4.15			
	MAHMA1001186	3.02	2.01	9.14	2.7	2.02	2.05	2.34	0.77	3.58			
35	MAHMA1001189	17.92	24.89	44.3	19.97	22.86	16.26	24.9	14.32	18.92			
	MAHMA1001191	10.11	15.9	19.2	7.11	13.06	8.79	15.3	10.81	16.52			
	MAHMA1001198	4.68	8.28	11.96	3.49	7.28	5.15	6.52	2.84	2.49			
	MAHMA1001202	30.38	37.04	84.46	54.02	29.63	36.02	18.16	17.9	40.13			
	MAHMA1001203	8.99	12.77	20.34	10.56	11.45	9.96	10.07	8.3	11.49			
	MAHMA1001206	14.27	17.3	21.28	14.44	9.47	13.95	15.09	7.44	16.29			
40	MAHMA1001208	3.37	1.85	7.28	0.99	1.62	0.23	1.58	0.63	2.48			
	MAHMA1001215	12.57	20.58	26.01	4.39	17.55	6.44	10.98	8.12	6.27			
	MAHMA1001220	12.2	14.97	20.97	18.99	11.55	15.4	10.87	5.54	12.65			
	MAHMA1001222	13.14	15.27	54.43	12	15.72	10.61	17.61	13.51	15.3			
	MAHMA1001223	5.47	11.04	23.87	8.51	8.76	8.32	9.09	3.98	9.79			
	MAHMA1001232	5.9	6.27	10.44	10.05	4.68	7.4	3.45	2.12	5.89			
45	MAHMA1001234	78.47	62.1	106.68	104.09	22.14	88.22	44.43	81.83	25.73			
	MAHMA1001237	4.23	3.61	4.51	2.39	3.8	1.85	3.83	2.61	1.63			
	MAHMA1001243	26.51	33.45	37.19	22.72	19.97	20.13	13.4	8.73	6.36	*	**	-
	MAHMA1001244	3.29	3.23	5.47	2.05	2.81	0.99	2.32	0.99	1.85			
	MAHMA1001249	32.97	38.57	66.31	31.72	40.81	37.57	44.31	25.2	27.49			
	MAHMA1001256	13.8	15.37	22.1	10.86	16.44	20.14	15.27	12.69	11.49			
50	MAHMA1001259	3.58	4.73	9.83	2.71	2.66	1.74	5.81	0.4	3.31			
	MAHMA1001260	15.61	19.65	19.14	11.02	7.29	22.35	9.03	6.24	10.39	**	-	
	MAHMA1001262	8.3	48.53	16.27	15.78	7.11	6.47	9.48	5.41	7.71			
	MAHMA1001268	7.26	6.65	12.6	5.74	6.05	4.12	3.69	3.37	4.04			
	MAHMA1001271	4.99	4.56	6.83	2.12	3.55	3.26	4.68	2.4	2.94	*	-	
	MAHMA1001274	197.38	283.08	348.26	54.68	91.98	70.74	33.35	19.31	25.39	*	**	-
55	MAHMA1001280	10.76	14.33	28.14	10.36	11.35	10.7	14.52	8.71	14.89			
	MAHMA1001283	4.11	3.31	10.89	0.34	3.46	2.07	2.94	1.3	1.97			
	MAHMA1001284	6.24	6.15	16.76	7	6.77	4.3	3.05	1.61	4.29			

Table 394

5	MAMMA1000824	52.75	82.46	54.98	22.75	86.37	46.11	53.27	21.38	20.22				
	MAMMA1000827	3.29	5.2	12.5	6.91	5.04	2.83	2.54	1.96	3.02				
	MAMMA1000831	31.08	35.42	39.96	20.36	22.38	25.87	17.52	21.92	26.79	*	*	-	-
	MAMMA1000838	4.58	6.92	8.85	2.9	2.71	3.02	3.13	2.4	1.8	*	*	-	-
	MAMMA1000839	4.7	5.1	9.24	1.82	2.67	4.2	2.02	1.34	1.48		*	-	-
	MAMMA1000841	4.43	2.78	9.96	2.71	2.83	2.07	3.01	0.81	2.24				
	MAMMA1000842	6.22	8.19	14.72	3.28	5.48	4.22	5.77	2.92	7.82				
10	MAMMA1000843	10.02	23.1	28.88	14.54	17.26	10.11	4.63	3.65	6.21	*		-	
	MAMMA1000845	2.57	2.89	10.34	0.88	3.81	2.04	1.66	1.53	3.4				
	MAMMA1000851	40.29	50.53	100.08	64.55	45.92	39.66	42.2	36.12	64.01				
	MAMMA1000854	3.86	5.86	8.12	2.91	1.76	3.56	1.11	0.53	1.24	*		-	
	MAMMA1000855	91.57	117.96	159.35	213.71	138.54	171.88	95.66	65.57	93.06				
	MAMMA1000856	3.13	2.69	5.73	1.9	1.05	2.66	1.82	0.46	1.81				
15	MAMMA1000859	3.59	5.65	10.37	1.76	1.83	1.2	3.38	1.09	1.24				
	MAMMA1000862	8.58	10.61	23.09	5.34	11.79	7.43	5.15	1.27	2.11				
	MAMMA1000863	17.14	20.47	45.55	14.36	24.31	16.09	24.81	12.68	16.34				
	MAMMA1000865	3.2	5.96	11.12	4.18	4.78	11.8	2.48	0	2.86				
	MAMMA1000867	12.55	18	37.97	18.03	16.36	15.97	9.25	8.53	14.8				
	MAMMA1000875	197.47	247.19	307.15	151.05	163.7	159.54	127.81	148.8	137.51	*	*	-	-
20	MAMMA1000876	4.29	6.54	6.75	2.84	2.75	3.43	2.51	1.58	2.61	*	*	-	-
	MAMMA1000877	2.84	3.61	4.96	2.05	1.13	1.68	2.93	0.77	2.2	*		-	-
	MAMMA1000878	5.22	9.31	15.62	6.72	6.36	5.06	10.1	4.12	5.11				
	MAMMA1000880	13.15	18.7	32.08	12.61	12.85	8.92	10.54	7.92	11.52				
	MAMMA1000881	6.97	11.85	26.23	10.06	11.84	7.67	8.29	6.16	7.32				
	MAMMA1000883	11.76	21.56	31.01	16.67	19.82	18.1	18.77	10.78	19.11				
25	MAMMA1000897	18.89	25.07	44.99	26.44	17.53	21.31	17.58	14.36	19.92				
	MAMMA1000898	4.54	4.74	8.26	5.15	4.11	4	3.9	1.83	1.2				
	MAMMA1000905	19.81	46.58	24.87	13.58	12.04	12.03	17.95	11.63	11.81				
	MAMMA1000906	13.7	9.74	18.2	5.83	6.94	4.39	4.39	2.34	1.82	*	*	-	-
	MAMMA1000908	8.3	13.83	18.11	10.1	14.32	5.63	5.09	3.21	3.69	*		-	-
	MAMMA1000911	13.12	16.04	26.46	13.81	12.11	7.57	12.29	5.96	12.25				
30	MAMMA1000914	2.41	3.47	11.09	1.11	2.25	1.29	1.59	0	0.87				
	MAMMA1000920	68.92	152.1	139.06	77.3	106.8	94.98	122.37	27.22	25.46				
	MAMMA1000921	3.68	5.93	12.67	7.11	6.12	6.97	3.45	2.44	2.92				
	MAMMA1000931	35.49	29.12	32.19	9.74	14.81	15.82	7.85	20.92	4.78	**	*	-	-
	MAMMA1000940	16.07	19.99	34.79	11.62	14.07	12.51	10.88	8.84	14.41				
	MAMMA1000941	36.94	28.9	55.32	14.85	46.69	25.1	44.15	31.17	20.09				
35	MAMMA1000942	19.99	18.82	58.05	28.3	25.6	20.38	16.91	16.31	24.51				
	MAMMA1000943	14.05	10.94	26.95	6.92	4.75	8.24	3.89	2.4	1.4	*		-	
	MAMMA1000952	3.98	5.7	16.6	3.37	4.97	3.51	7.54	1.89	3				
	MAMMA1000956	5.62	5.18	14.83	6.99	6.03	6.18	4.85	2.51	6.15				
	MAMMA1000957	0.61	0.47	2.13	0.38	0.27	1.52	0.16	0	0				
	MAMMA1000962	26.82	25.6	21.85	10.69	8.37	7.64	7.43	8.92	3.07	**	**	-	-
40	MAMMA1000966	13.96	15.32	18.62	3.18	4.77	5.42	5.48	5.55	2.07	**	**	-	-
	MAMMA1000968	3.82	3.17	8.04	2.19	4.04	1.69	2.56	0.98	1.54				
	MAMMA1000972	109.05	38.4	166	38.59	38.3	60.84	51.28	65.81	23.12				
	MAMMA1000973	4.24	3.6	8.43	1.16	3.43	2.39	3.18	1.23	1.88				
	MAMMA1000975	4.83	5.04	16.94	6.44	6.12	4.99	5.27	1.93	5.12				
	MAMMA1000976	10.4	12.69	31.13	7.88	10.7	11.37	8.68	4.42	4.88				
45	MAMMA1000979	15.33	18.37	49.49	29.29	22.41	26.44	15.03	8.9	16				
	MAMMA1000986	11.83	20.17	19.1	14.27	9.07	14.47	9.64	5.8	3.91	*		-	
	MAMMA1000987	35.85	39.55	84.06	67.43	62.8	61.8	46.73	25.89	38.89				
	MAMMA1000988	12.53	13.76	42.26	22.92	17.53	17.38	12.36	7.34	12.64				
	MAMMA1000994	6.93	10.09	26.01	16.14	13.54	12.99	6.56	3.79	6.83				
	MAMMA1000996	3.85	8.09	10.95	3.78	4.4	5.85	1.97	0.73	2.47	*		-	
50	MAMMA1001003	11.87	14.9	34.42	24.16	21.08	20.65	13.23	10.02	16.87				
	MAMMA1001007	4	4.13	6.2	3.14	6.73	3.28	4.14	1.49	4.79				
	MAMMA1001008	20.12	23.76	44.26	26.75	20.79	27.13	18.57	19.08	24.22				
	MAMMA1001013	9.3	12.99	14.08	13.97	3.4	7.56	6.85	5.46	6.5	*		-	
	MAMMA1001014	3.74	2.89	7.41	3.98	5.49	5.98	4.67	4.19	1.31				
	MAMMA1001021	58.82	76.19	93.74	67.98	54.85	51.75	72.35	45.95	58.5				
55	MAMMA1001024	88.7	92.15	131.2	139.16	94.51	99.4	87.98	68.05	106.24				
	MAMMA1001025	69.9	102.4	41.66	70.13	53.68	30.76	22.6	25.23	39.06				
	MAMMA1001026	80.18	111.56	110	73.96	84.44	72.45	97.15	91.69	100.75				

Table 393

	MAMMA1000567	2.83	9.61	12.53	4.59	4.66	3.08	3.91	1.78	2.49			
	MAMMA1000576	14.4	23.94	41.43	6.77	7.53	16.66	14.73	20.93	24.41			
5	MAMMA1000582	342.62	404.06	589.09	206.24	397.1	307.2	196.5	234.99	295.12			
	MAMMA1000583	3.44	4.94	9.73	2.95	5.37	2.97	1.64	1.64	2.31			
	MAMMA1000585	10.36	9.76	14.3	7.6	5.3	4.54	9.1	4.55	7.57	*	-	
	MAMMA1000587	9.1	16.85	23.22	12.86	13.24	10.32	9.64	5.62	11			
	MAMMA1000591	24.67	34.19	52.67	17.16	22.85	23.65	15.05	11.26	16.03			
10	MAMMA1000594	25.66	32.59	51.71	40.6	49.12	36.4	18.51	9.64	20.88			
	MAMMA1000597	10.5	17.98	38.05	21.57	17.46	15.09	12.1	5.53	10.65			
	MAMMA1000605	8.03	8.9	14.02	3.22	4.68	8.25	3.07	3.06	5.3	*	-	
	MAMMA1000612	3.76	3.67	7.3	1.87	2.17	1.53	2.62	0.72	2.53			
	MAMMA1000614	8.91	9.36	17.48	7.74	5.34	6.31	7.11	2.55	5.61			
	MAMMA1000616	10.09	7	22.98	5.69	13.58	6.11	4.8	4.98	3.1			
15	MAMMA1000621	18.37	26.27	42.33	24.96	17.41	18.13	16.86	13.76	19.28			
	MAMMA1000623	5.51	4.72	11.4	2.5	4.83	2.59	2.52	1.82	1.65			
	MAMMA1000625	3.19	6.18	12.95	3.95	4.29	3.37	3.1	1.01	2.69			
	MAMMA1000635	3.08	4.28	9.79	5.68	3.5	2.91	2.33	0.32	2.12			
	MAMMA1000643	3.16	4.8	8.96	4.43	4.19	14.5	3.27	1.73	4.43			
	MAMMA1000646	53.05	63.8	128.68	68.06	63.84	76.26	48.7	34.2	32.4			
20	MAMMA1000652	7.49	6.59	12.24	7.34	7.8	10.34	5.92	2.92	1.26			
	MAMMA1000657	4.53	4.12	7.44	5.71	2.66	4.08	1.96	2.59	1.6	*	-	
	MAMMA1000664	18.71	22.51	38.34	17.17	14.96	16	10.64	12.09	11.81			
	MAMMA1000667	8.69	17.36	17.2	3.58	6.25	8.92	3.77	5.36	2.7	*	-	
	MAMMA1000668	23.14	46.99	61.08	27.38	33.9	32.87	41.85	23.92	33.78			
	MAMMA1000669	3.03	3.5	9.1	2.2	6.51	1.81	2.38	2.13	1.46			
25	MAMMA1000670	4.89	6.68	9.53	5.24	3.91	6.43	9.4	3.5	1.93			
	MAMMA1000672	127	118.39	232.08	362.26	175.38	245.8	112.44	89.34	106.16	*	**	- -
	MAMMA1000681	42.17	56.4	66.57	28.1	10.46	18.45	7.15	8.3	5.57	*	**	- -
	MAMMA1000684	6.35	5.06	11.76	8.34	4.22	4.13	4.18	2.86	3.12			
	MAMMA1000696	7.53	9.04	16.29	12.54	11.44	15.76	4.16	3.21	5.89			
	MAMMA1000702	2.37	2.75	7.2	4.27	3.94	4.85	2.38	1.14	1.76			
30	MAMMA1000706	7.11	13.17	13.46	6.85	13.62	11.39	5.11	2.82	4.43	*	-	
	MAMMA1000707	5.79	9.88	8.46	5.04	7.37	7.72	4.37	2.73	5.56	*	-	
	MAMMA1000713	13.18	25.83	23.83	10.07	8.95	6.62	3.02	4.1	4.74	*	*	- -
	MAMMA1000714	21.05	29.79	38.76	24.92	18.64	21.26	20.38	21.53	20.97			
	MAMMA1000718	7.11	5.26	13.51	4.07	3.72	3	2.71	1.31	2.37			
	MAMMA1000720	56.24	56.67	100.19	108.88	65.38	82.83	53.41	37.73	50.75			
35	MAMMA1000723	83.95	62.78	122.37	158.83	112.68	140.59	85.25	48.89	63.98			
	MAMMA1000731	13.21	20.89	24.92	19.74	19.6	18.49	13.07	8.64	14.74			
	MAMMA1000732	4.13	6	9.07	5.99	9.97	7.96	4.16	1.33	3.64			
	MAMMA1000733	12.22	18.56	20.77	15.57	15.03	11.37	17.6	9.79	15.79			
	MAMMA1000734	4.23	6.43	10.91	7.98	3.73	6.91	3.13	3.72	3.43			
	MAMMA1000736	4.58	4.07	5.19	5.1	2.18	3.84	1.49	1.8	2.72	**	-	
40	MAMMA1000738	27.47	31.39	43.2	35.52	27.09	30.72	11.08	15.85	19.91	*	-	
	MAMMA1000744	11.43	16.18	20.32	9.63	9.84	10.4	16.64	15.89	17.27			
	MAMMA1000746	5.14	6.4	11.37	3.9	6.11	4.85	2.11	2.43	3.5			
	MAMMA1000748	2.95	3.98	9.11	2.66	4.26	3.02	2.63	1.16	2.48			
	MAMMA1000751	3.53	4.94	7.73	2.22	4.21	3.15	2.53	1.56	2.63			
	MAMMA1000752	9.18	9.76	16.46	16.7	13	12.37	10.17	5.29	10.03			
45	MAMMA1000757	17.77	21.89	38.71	25.58	19.94	21.44	12.48	15.54	20.02			
	MAMMA1000760	6.79	10.56	9.64	3.73	4.03	3.96	1.97	4.15	4.68	*	*	- -
	MAMMA1000761	4.06	4.2	7.68	1.53	2.88	3.67	1.98	1.33	1.45	*	-	
	MAMMA1000775	4.12	5.64	14.4	7.07	2.94	5.07	2.8	2.71	4.1	*	-	
	MAMMA1000776	26.15	27.77	36.92	48.47	28.15	33.74	22.66	12.7	18.46	*	-	
	MAMMA1000778	38.93	73.17	60.3	15.58	59.91	40.66	15.6	19.11	12.75	*	-	
50	MAMMA1000781	10.71	16.72	21.48	6.1	15.65	13.14	10.25	9.83	6.19			
	MAMMA1000782	5.28	6.96	8.31	4.59	9.33	5.12	3.95	1.7	2.78	*	-	
	MAMMA1000784	12.93	21.78	27.25	9.87	8.6	8.24	5.41	7.95	18.83	*	-	
	MAMMA1000786	14.83	19.2	21.42	14.77	17.9	16.09	9.03	8.51	18.38			
	MAMMA1000798	10.63	12.91	13.99	6.66	4.6	7.92	6.32	3.01	7.59	*	*	- -
	MAMMA1000802	10.59	10.38	19.17	11.33	9.98	8.79	5.57	3.43	5.21	*	*	- -
55	MAMMA1000810	6.03	6.18	13.72	6.5	6.85	7.38	2.82	2.22	3.59			
	MAMMA1000813	22.11	37.42	37.78	16.99	20.99	20.43	27.71	18.27	23.96			
	MAMMA1000814	1.64	5.26	13.69	2.27	7.88	1.6	3.8	1.06	1.8			

Table 392

	MAMMA1000305	32.01	34.24	58.88	37.52	42.02	39.78	30.9	18.25	28.49			
	MAMMA1000307	7.32	11.17	15.9	9.52	5.29	9	4.83	4.75	5.32			
5	MAMMA1000309	13.21	12.45	16.61	13.37	8.5	8.68	5.45	4.69	8.26	**	-	
	MAMMA1000312	3.8	4.66	7.38	5.51	4.44	4.58	3.65	2.16	2.53			
	MAMMA1000313	5.23	3.78	8.22	6.32	8.18	4.37	4.06	1.99	2.36			
	MAMMA1000331	3.57	2.97	7.78	3.29	3.56	6.01	3.84	1.96	2.66			
	MAMMA1000335	1.93	3.97	7.5	5.14	3.16	3.87	4.24	1.18	2.51			
10	MAMMA1000339	4.85	7.71	8.76	4.7	9.96	8.63	2.97	1.94	3.31	*	-	
	MAMMA1000340	13.8	21.06	20.08	19.65	20.75	20	20.39	11.88	8.29			
	MAMMA1000348	6.6	5.86	10.24	5.86	4.39	5.28	1.91	2.51	3.44	*	-	
	MAMMA1000356	48.67	74.66	87.35	53.83	37.37	46.85	35.71	39.77	39.98			
	MAMMA1000358	9.54	13.7	18.74	8.76	9.05	7.26	6.27	7.75	6.04			
	MAMMA1000360	4.42	8.35	14.28	5.25	6.74	5.72	3.05	4.95	4.76			
15	MAMMA1000361	62.5	108.93	116.66	101.62	117.16	105.86	87.17	73.39	114.81			
	MAMMA1000363	4.98	6.13	8.18	4.8	3.56	6.25	3.82	2.21	3.44	*	-	
	MAMMA1000370	5.17	12.45	10.14	5.35	7.18	7.58	3.49	1.29	2.83	*	-	
	MAMMA1000371	2.39	4.39	9.15	3.68	6.58	4.68	2.76	1.36	2.71			
	MAMMA1000372	17.51	21.43	34.07	12.37	14.77	13.39	11.25	17.12	32.9			
	MAMMA1000385	2.59	3.13	9.36	4.39	3.21	2.85	3.43	3.75	2.78			
20	MAMMA1000388	15.23	16.36	27.37	8.49	18.09	20.18	11.94	20.32	17.28			
	MAMMA1000395	2.71	-2.65	9.67	2.23	1.16	2.27	1.54	2.51	1.57			
	MAMMA1000402	17.85	14.83	40.38	21.39	34.95	27.07	11.67	5.88	6.61			
	MAMMA1000403	4.76	5.81	8.95	2.66	5.93	4.73	3.53	2.03	3.28			
	MAMMA1000410	8.38	10.39	14.35	8.64	10.52	7.1	6.79	5.15	5.87	*	-	
	MAMMA1000413	13.08	31.41	34.59	25.16	26.53	19.73	20.54	9.59	10.58			
25	MAMMA1000414	26.2	35.15	53.64	41.13	33.28	34.97	22.92	16.5	31.45			
	MAMMA1000416	93.85	138.41	125.21	46.14	53.85	91.17	35.64	64.76	53.2	*	*	-
	MAMMA1000421	12.39	14.43	18.48	7.65	12.15	12.39	5.74	4.49	4.48	**	-	-
	MAMMA1000422	142.54	131.07	148.31	147.75	137.43	173.26	122.42	112.28	99.77	*	-	-
	MAMMA1000423	5.22	2.76	10.5	5.24	3.96	3.14	2.65	0.29	3.52			
	MAMMA1000424	16.73	20.92	35.59	20.88	21.28	23.97	15.73	11.76	16.32			
30	MAMMA1000429	9.36	15.84	18.54	7.22	10.71	6.57	4.21	3.75	2.1	*	-	
	MAMMA1000431	1.71	4.16	6.38	1.45	4.13	3.04	2.28	2.32	2.5			
	MAMMA1000432	63.4	133.96	122.67	9.74	51.75	20.93	35.92	64.94	54.93	*	-	
	MAMMA1000437	7.75	15.03	14.39	12.33	7.79	9.82	5.47	8.03	11.07			
	MAMMA1000444	7.3	9.64	12.3	7.17	6.74	4.04	4.84	3.63	6.39	*	-	
	MAMMA1000446	72.49	65.48	72.84	97.18	57.65	80.67	67.4	46.57	58.28			
35	MAMMA1000449	18.1	22.35	31.63	13.49	17.76	14	17.72	12.22	18.97			
	MAMMA1000457	18.4	25.87	17.2	2.76	17.71	11.5	15.1	3.75	16.53			
	MAMMA1000458	7.78	16.72	16.75	9.69	10.94	4.65	4.37	2.84	7.72			
	MAMMA1000468	16.03	28.35	43.07	20.2	27.28	30.25	21.32	9.68	14.64			
	MAMMA1000472	6.66	14.63	17.94	8.59	8.34	7.38	4.92	4.41	8.54			
	MAMMA1000473	18.92	28.23	28.07	15.83	15.65	17.15	17.93	13.07	20.76	*	-	
40	MAMMA1000477	3.29	5.76	7.67	2.77	2.69	3.84	2.99	1.75	1.93			
	MAMMA1000478	3.8	4.66	7.21	2.84	2.45	2.89	2.43	1.21	0.87	*	-	
	MAMMA1000483	141.82	256.33	267.79	98.09	257.9	180.98	169.43	158.7	175.76			
	MAMMA1000490	11.74	16.92	30.01	11.37	13.37	11.4	15.98	7.97	17.05			
	MAMMA1000495	29.48	25.33	53.38	30.03	31.05	22.35	19.18	13.04	15.93			
	MAMMA1000500	4.21	8.78	19.16	2.1	8.22	5.73	7.01	0.97	6.78			
45	MAMMA1000501	3.3	3.92	7.8	5.4	3.37	3.89	2.1	0.03	4.12			
	MAMMA1000503	8.55	8.62	11.61	8.76	3.91	6.21	3.65	2.45	1.88	**	-	
	MAMMA1000506	19.56	37.71	40.79	41.28	34.62	14.86	17.29	25.75	7.33			
	MAMMA1000510	6.43	9.68	12.68	8.8	8.42	5.47	7.38	4.77	5.1			
	MAMMA1000515	16.44	20.18	30.29	25.14	20.54	20.81	15.29	9.87	15.02			
	MAMMA1000516	5.05	38.96	14.92	3.26	5.69	7.88	10.33	1.91	1.96			
50	MAMMA1000522	27.99	43.17	68.38	38.83	18.51	22.66	24.89	20.61	23			
	MAMMA1000524	5.97	11.73	10.41	4.38	10.1	5.92	5.73	2.46	2.76	*	-	
	MAMMA1000528	3.99	4.53	7.16	4.83	4.31	6.82	4.96	1.75	1.88			
	MAMMA1000534	20.75	20.93	19.83	14.18	15.4	10.89	14.95	12.85	11.39	**	**	-
	MAMMA1000541	7.6	8.83	11.36	1.44	4.17	2.11	2.2	1.84	1.98	**	**	-
	MAMMA1000550	2.63	1.99	8.54	2.03	3.94	1.32	3.32	0.71	2.83			
55	MAMMA1000556	761.19	1381.9	1153	543.64	1435.2	1032.9	1229.3	435.34	855.29			
	MAMMA1000559	6.09	5.3	11.69	2.67	3.19	2.37	2.27	0.81	3.09			
	MAMMA1000565	3.15	5.75	13.35	1.46	4.28	0.82	2.18	0.1	1.13			

Table 391

	MAMMA1000025	3.49	6.73	9.73	4.76	8.15	5.09	3.99	1.9	3.02			
5	MAMMA1000043	8.59	8.02	15.22	6.68	3.56	4.76	2.11	1.9	6.1			
	MAMMA1000045	9.03	8.98	17.21	11.43	6.71	9.34	5.16	4.47	6.29			
	MAMMA1000046	41.47	53.57	56.97	30.34	27.99	31.56	37.45	30.61	48.64	*		
	MAMMA1000055	10.6	18.89	12.32	4.6	6.26	5.89	3.62	4.44	2.53	*	*	-
	MAMMA1000057	3.76	2.41	12.19	2.23	1.8	1.43	1.21	1.53	2.81			
	MAMMA1000060	10.21	8.2	18.54	5.99	7.11	5.88	5.98	4.17	6.29			
10	MAMMA1000069	143.75	270.15	259.45	49.26	67.48	38.63	33.36	39.16	49.31	*	*	-
	MAMMA1000084	3.34	5.16	10.93	3.47	4.79	2.05	3	0.67	2.88			
	MAMMA1000085	3.86	4.9	7.86	4.65	4.18	4.46	2.36	1.98	8.94			
	MAMMA1000092	1.88	4.62	8.98	3.96	2.75	3.09	2.27	0.07	1.18			
	MAMMA1000096	3.52	3.48	7.14	2.58	2.53	2.5	0.67	1.83	1.35			
	MAMMA1000097	4.98	3.73	8.88	2.04	2.02	1.49	1.93	0.71	1.64	*		-
15	MAMMA1000102	16.31	20.65	29.78	15.97	14.1	11.62	28.48	21.84	17.26			
	MAMMA1000103	24.25	30.87	39.49	18.45	28.7	17.72	21.24	13.56	7.51	*		-
	MAMMA1000106	8.95	10.76	22.4	12.82	11.13	10.63	6.32	4.47	6.45			
	MAMMA1000117	2.83	6.06	12.12	2.65	4.11	2.11	5.97	1.27	1.18			
	MAMMA1000118	8.06	8.77	19.58	14.56	12.82	9.23	5.17	8.7	14.06			
	MAMMA1000129	4	2.76	5.79	3.46	3.6	1.79	1.7	2.01	2.69			
20	MAMMA1000133	5.05	7.04	6.39	3.71	2.31	3.3	8.78	9.22	3.38	*		-
	MAMMA1000134	7.25	6.71	12.23	3.97	5.46	2.89	5.9	4.64	5.77			
	MAMMA1000139	8.81	13.92	27.94	9.57	7.21	7.44	10.8	7.32	10.69			
	MAMMA1000141	10.59	12.38	22.86	16.46	15.91	12.46	8.35	6.81	8.98			
	MAMMA1000143	47.46	88.55	73.36	60.42	67.69	52.39	70.22	26.61	57.3			
	MAMMA1000150	7.57	5.6	14.19	2.79	8.22	2.73	5.6	1.58	3.48			
25	MAMMA1000155	11.41	12.76	22.27	10.11	6.62	6.85	2.76	1.55	4.85	*		-
	MAMMA1000163	15.02	19.25	32.92	19.25	12.39	13.68	10.19	9.3	14.14			
	MAMMA1000171	7.6	10.37	13.24	2.98	2.32	1.89	2.54	1.28	1.33	**	**	-
	MAMMA1000173	6.75	5.29	9.57	2.32	2.23	1.04	2.07	1.37	0.99	*	*	-
	MAMMA1000175	3.46	6.59	11.02	3.52	2.3	2.43	3.53	1.51	2.66			
	MAMMA1000183	25.33	42.54	44.08	23.65	35.89	24.05	16.46	12.47	14.16	*		-
30	MAMMA1000191	5.92	8.95	18.7	5.23	6.43	6.14	3.97	2.29	6.56			
	MAMMA1000192	11.99	14.56	36.16	26.22	18.77	12.98	9.82	7.53	11.64			
	MAMMA1000193	3.19	3.37	6.86	4.95	4.82	4.54	1.06	1.39	2.14			
	MAMMA1000198	47.48	56.45	65.4	32.82	40.48	48.35	28.71	22.96	42.97	*		-
	MAMMA1000204	11.23	12.69	25.02	6.96	11.44	9.78	5.53	7.76	6.1			
	MAMMA1000207	12.4	7.47	19.55	6.26	5.09	4.84	6.03	8.39	7.97			
35	MAMMA1000214	8.42	8.75	15.08	9.58	9.32	7.56	7.33	4.95	7.33			
	MAMMA1000220	22.52	15.12	36.99	5.02	6.29	6.82	6.38	3.87	3.09	*	*	-
	MAMMA1000221	10.4	16.64	24.93	5.03	11.07	6.49	6.69	3.21	1.94	*		-
	MAMMA1000226	13.64	11.97	28.4	5.25	12.91	9.41	26	12.92	3.5			
	MAMMA1000227	14.38	17.15	40.21	23.86	16.67	19.14	13.63	7.29	12.46			
	MAMMA1000230	3.53	5.09	7.34	3.64	2.92	4.31	1.76	1.24	2.65	*		-
40	MAMMA1000241	2.76	3.88	6.29	2.63	1.66	1.65	1.27	0.15	1.78	*		-
	MAMMA1000245	6.45	7.89	22.51	9.95	8.51	9.68	6.76	3.1	4.13			
	MAMMA1000248	21.38	25.61	55.84	30.22	20.66	19.73	24.19	11.24	13.89			
	MAMMA1000251	5.31	4.84	10.73	2.94	4.44	3.42	3.67	1.53	2.11			
	MAMMA1000254	2.32	3.23	8.55	2.43	2.71	1.01	0.9	0.97	0.54			
	MAMMA1000257	2.97	3.77	9.11	1.37	3.58	0.82	2.27	0.96	1.23			
45	MAMMA1000262	45.74	36.4	0	0.39	0	0	34.58	23.42	46.99			
	MAMMA1000264	9.34	12.72	36.1	26.8	15.94	23	7.31	7.08	9.25			
	MAMMA1000266	0.66	1	3.04	2.63	2.61	0.59	0.15	0	1			
	MAMMA1000270	13.12	14.9	18.21	14.67	11.46	11.77	8.4	9.61	6.99	*		-
	MAMMA1000271	3.88	3.72	7.35	4.08	3.89	3.79	1.47	2.83	1.69			
	MAMMA1000277	41.4	51.52	103.66	30.74	33.8	53.69	17.12	13.52	16.27			
50	MAMMA1000278	11.35	16.91	37.8	22.57	16.53	16.11	10.35	6.71	12.84			
	MAMMA1000279	3.26	3.91	6.49	3.15	3.6	2.86	2.52	0.56	1.77			
	MAMMA1000283	29.38	33.95	49.97	69.77	34.21	54.3	27.86	21.79	21.83			
	MAMMA1000284	5.55	5.62	6.77	4.76	4.06	5.69	4.18	2.05	4.21	*		-
	MAMMA1000287	18.44	27.31	36.41	15.51	4.83	9.89	4.34	3.99	2.59	*	*	-
	MAMMA1000294	14.59	17.3	29.88	19.53	15.7	20.07	17.98	12.1	15.45			
55	MAMMA1000298	4.06	4.13	7.44	4.06	5.42	8.4	4.25	2.39	4.19			
	MAMMA1000302	3.99	3.48	11.09	6.9	6.88	5.91	4.93	2.1	3.71			
	MAMMA1000303	1.77	4.16	5.21	2.14	3.15	3.07	3.83	0.6	1.85			

Table 390

	HEMB81002635	3.81	5	11.37	1.21	3.62	1.08	2	1.05	2.82			
5	HEMB81002644	6.38	7.38	19.6	6.87	6.62	5.91	7.04	3.57	7.83			
	HEMB81002654	44.15	47.27	98.13	55.06	50.78	51.15	27.74	20.25	41.98			
	HEMB81002661	16.29	22.35	33.51	18.43	18.3	14.62	15.42	10.61	19.12			
	HEMB81002663	12.29	20.14	24.54	5.77	16.48	12.72	3.66	7.87	3.2		*	-
	HEMB81002664	26.04	27.32	31.97	12.71	14.38	12.28	15.87	10.5	16	**	**	-
	HEMB81002677	2.85	4.67	7.35	1.19	3.8	1.22	2.58	0.54	2.04			
10	HEMB81002683	408.12	530.85	581.04	161.92	536.41	322.12	493.61	190.86	318.34			
	HEMB81002684	5.89	14.15	21.74	5.64	5.67	5.5	2.78	0.53	2.33			
	HEMB81002686	2.08	2.21	13.73	0.94	2.87	1.96	2.53	0.03	2.4			
	HEMB81002692	6.27	8.28	18.48	7.56	4.81	4.6	4.23	1.22	6.95			
	HEMB81002693	37.13	59.89	50.13	8.18	9.08	32.77	12.89	24.23	25.51	*	*	-
	HEMB81002697	12.19	19.73	29.39	11.74	11.32	12.96	13.01	7.86	10.37			
15	HEMB81002699	12.29	7.97	29.87	3.99	12.05	3.69	6.19	2.34	1.91			
	HEMB81002702	4.21	4.82	9.81	4.02	2.81	2.59	2.32	3.46	2.32			
	HEMB81002705	42.01	58.09	92.41	42.52	47.93	35.39	82.62	52.43	59.52			
	HEMB81002712	82.07	113.22	149.92	77.66	93.78	72.72	96.78	72.75	86.71			
	IMR321000028	2.68	4.75	13.62	4.72	4.88	2.62	2.74	1.07	1.92			
20	IMR321000031	7.53	10.04	0	0.26	16.99	0.79	7.55	3.43	9.54			
	IMR321000034	9.42	10.78	32.44	15.73	16.03	17.46	6.96	5.24	7.94			
	IMR321000039	1.72	2	2.81	3.07	3.78	1.79	0.76	0	1.77			
	IMR321000044	5.65	7.08	18.45	11.11	7.14	7.03	4.83	2.08	3.62			
	IMR321000063	10.51	11.67	19.12	13.18	11.6	12.05	6.47	7.76	7.56			
	IMR321000085	3.33	5.05	8.66	2.61	3.29	3.45	1.96	0.92	3.07			
	IMR321000089	3.15	5.89	14.29	4.37	6.53	5.77	3.28	1.07	2.45			
25	IMR321000091	59.8	78.47	131.41	171.62	125.26	143.68	43.76	27.21	45.98			
	LIVER1000004	4.13	4.93	5.75	3.08	2.76	4.02	2.18	1.07	1.36	**		-
	LIVER1000008	10.35	13.6	22.11	11.7	10.53	14.7	11.26	5.83	10.54			
	LIVER1000011	59.6	50.92	129.92	124.27	66.83	94.45	48.23	37.68	48.06			
	LIVER1000022	2.42	3.87	7.13	5.18	2.89	5.96	3.89	0.85	2.2			
30	LIVER1000025	11.19	21.41	14.85	14.92	17.52	18.14	8.19	3.93	7.48	*		-
	LIVER1000030	5.79	4.25	9.83	9.98	5.67	5.82	4.77	2.89	3.63			
	LIVER1000045	4.77	5.86	5.47	6.22	7.71	6.61	4.81	3.96	4.97			
	LIVER1000046	23.39	30.93	40.35	28.67	14.67	17.18	8.63	19.03	18.73			
	LIVER1000072	9.51	20.72	24.34	17.51	14.67	15.78	8.13	9.06	13.17			
	LIVER1000077	7.51	7.91	13.85	13.48	8.07	9.05	5.23	4.4	6.07			
35	LIVER1000080	5.07	5.78	10.96	6.47	4.34	6.39	3.86	2.15	4.56			
	LIVER1000086	11.03	13.61	23.46	8.67	23.05	12.3	11.46	8.1	16.78			
	LIVER1000092	10.59	15.75	23.53	19.18	22.61	13.71	8.85	6.89	7.35			
	LIVER1000095	51.53	68.53	98.85	69.98	49.46	69.44	74.79	43.47	64.56			
	LIVER1000097	8.95	15.6	14.45	10.75	13.82	9.14	5.8	2.85	3.46	*		-
	LIVER1000098	41.49	40.97	103.47	50.66	24.2	43.26	26.16	32.27	27.07			
40	LIVER1000100	3.55	4.13	7.44	3.5	3.68	3.78	2.58	3.54	3.3			
	LIVER1000101	44.76	62.72	56.32	36.92	31.17	34.9	40.25	36.83	38.75	*	*	-
	LIVER1000106	3.93	4.77	8.45	2.61	3.61	4.62	2.14	3.23	3.22			
	LIVER1000108	6.19	8.29	14.94	4.01	4	6.54	5.47	6.84	5.37			
	LIVER1000115	1.31	3.43	5.31	1.07	1.91	1.17	0.78	1.21	1.27			
	LIVER1000120	5.87	8	12.35	7.59	6.45	6.85	5.23	2.59	6.22			
45	LIVER1000138	5.5	7.3	19.82	7.01	10.38	6.37	5.23	4.12	9.23			
	LIVER1000146	4.13	5.06	8.45	8.15	4.26	4.91	1.55	2.41	3.37			
	LIVER1000148	10.81	16.45	14.93	9.88	11.05	13.74	6.11	8.29	5.5	*		-
	LIVER1000157	4.51	3.83	6.67	3.38	5.06	4.69	1.48	2.24	4.29			
	LIVER1000161	85.1	101.44	127.49	155.4	95.91	123.47	63.19	53.6	81.33			
	LIVER1000167	9.34	7.8	16.08	6.28	7.39	6.18	4.07	3.11	4.06	*		-
50	LIVER1000174	14.38	23.84	29.19	10.08	10.54	10.6	4.27	1.62	4.26	*	*	-
	LIVER1000185	15.55	11.69	24.81	3.86	12.76	7.24	6.4	3.76	4.58	*	*	-
	LIVER1000187	40.28	72.32	62.92	52.19	53.14	45.8	30.48	15.14	10.77	*	*	-
	LIVER1000190	13.01	13.24	16.57	5.85	5.19	5.07	2.85	4.77	5.27	**	**	-
	LIVER1000192	3.52	6.68	4.51	4.92	4.75	3.34	0.75	1.7	3.55			
	MANNA1000009	5.18	9.12	9.31	4.2	4.16	3.51	1.94	1.4	2.13	*	*	-
	MANNA1000015	5.78	4.65	11.74	5.4	3.83	5.36	2.48	1.65	3.15			
55	MANNA1000019	8.71	5.86	17.78	9.05	5.32	5.22	3.67	4.83	8.34			
	MANNA1000020	101.37	73.62	112.56	134.32	82.42	106.51	82.61	44.53	81.2			
	MANNA1000024	19.83	36.9	36.17	7.46	24.94	15.61	12.77	15.51	6.89	*		-

Table 389

	HEM881002326	6.79	10.97	11.32	21.02	12.83	6.76	6.12	3.86	6.53		
5	HEM881002327	29.86	39.37	46.64	28.75	34.27	35.49	23.76	23.05	28.43		
	HEM881002329	12.21	17.09	28.97	18.73	8.27	19.71	9.42	11.59	11.1		
	HEM881002340	3.57	5.21	5.98	7.77	3.22	9.25	5.09	2.16	1.79		
	HEM881002342	6.21	7.8	10.28	6	3.29	3.42	3.33	3.85	5.15	*	-
	HEM881002358	32.09	38.53	65.28	39.45	32.86	34.65	33.44	21.67	27.85		
	HEM881002359	3.96	3.69	6.45	3.93	4.64	4.39	3.95	0.63	2.74		
10	HEM881002364	3.43	4.07	8.33	6.97	4.3	5.26	4.27	1.48	3.01		
	HEM881002366	7.13	10.16	14.56	9.95	9.83	9.89	9.48	6.8	10.03		
	HEM881002371	17.53	29.76	21.32	21.87	21.71	16.54	9.41	9.09	4.87	*	-
	HEM881002381	7.92	12.33	16.15	8.95	3.96	8.62	4.76	5.05	4.4	*	-
	HEM881002383	135.68	179.56	228.28	286.57	191.47	226.83	107.86	116.89	120.62		
	HEM881002387	23.79	23.11	43.55	20.25	12.08	15.04	3.85	6.21	4.42	*	-
15	HEM881002409	13.31	12.51	22.34	8.77	11.21	10.89	4.44	4.37	4.87	*	-
	HEM881002413	3.49	3.12	9.08	4.81	2.78	4	1.3	1.86	1.66		
	HEM881002415	2.64	3.61	7.39	2.66	2.05	3.42	0.86	1.28	2.41		
	HEM881002424	6.07	9.92	9.97	6.37	9	12.89	6.72	4.73	6.15		
	HEM881002425	4.48	7.81	11.5	5.69	9.56	5.52	4.13	1.79	5.11		
	HEM881002427	29.25	30.49	60.22	43.33	23.55	33.03	24.65	22.4	31.01		
20	HEM881002442	6.2	17.54	17.42	9.79	7.28	9.01	4.36	12.11	11.81		
	HEM881002447	110.71	207.61	155.4	83.72	84.2	96.3	26.65	37.71	31.85	*	-
	HEM881002453	5.36	5.98	9.95	2.72	2.55	3.56	2.41	1.81	3.86	*	-
	HEM881002457	16.89	16.27	19.17	6.78	17.09	11.88	7.07	5.54	6.6	**	-
	HEM881002458	23.11	31.15	52.26	31.43	27.93	29.52	25.75	14.57	26.97		
	HEM881002463	24.77	42.25	57.37	32.51	37.48	24.85	28.18	20.41	30.3		
25	HEM881002465	3.27	5.17	9.01	3.69	4.27	4.31	3.51	0.9	2.26		
	HEM881002477	4.25	7.51	5.14	7.06	3.76	4.24	1.98	3.33	2.58	*	-
	HEM881002479	20.58	32.93	33.42	17.09	21.13	16.76	11.95	15.32	21.36		
	HEM881002489	48.69	62.6	67.58	34.04	44.48	57.32	35.3	36.02	39.44	*	-
	HEM881002492	11.53	17.66	26.19	13.16	10.94	12.55	11.2	5.85	15.28		
	HEM881002495	5.18	3.21	9.5	2.92	4.34	3.1	3.82	2.25	3.88		
30	HEM881002502	34.71	31.95	38.03	46.31	31.48	35.99	26.54	23.82	25.15	**	-
	HEM881002509	11.07	15.55	26.84	12.73	16.51	11.29	10.77	7	12.74		
	HEM881002510	28.51	45.37	36.84	31.94	38.1	20.25	23.53	21.31	17.08	*	-
	HEM881002520	14.34	15.62	18.28	10.3	7.17	4.37	1.27	2.73	3.96	*	**
	HEM881002522	27.23	55.84	54.45	41.8	39.8	28.31	15.32	22.78	36.38		
	HEM881002527	2.56	2.91	5.88	1.86	2.03	3.34	1.71	1.04	2.87		
35	HEM881002530	5.16	6.33	11.67	5.25	3.42	5.09	4.28	2.92	7.5		
	HEM881002531	26.73	31.41	45.92	22.42	21.99	29.07	24.79	13.7	18.84		
	HEM881002534	27.18	46.28	57.65	25.13	29.76	23.09	34.17	26.17	36.08		
	HEM881002536	13.04	20.55	23.13	9.22	13.07	13	15.33	12.24	18.4		
	HEM881002544	23	32.2	43.29	21.03	30.23	27.52	19.17	15.09	14.85		
	HEM881002545	3.59	4.33	9.7	5.81	3	3.66	1.12	0.96	13.79		
40	HEM881002550	7.33	13.21	13.27	3.83	6.5	4.73	2.81	3.61	2.61	*	*
	HEM881002556	19.36	24.38	22.91	7.86	11.05	11.87	7.3	3.24	5.41	**	**
	HEM881002571	7.32	4.72	8.5	4.01	3.71	2.33	1.87	1.16	2.04	*	*
	HEM881002579	3.21	4.8	10.5	2.3	2.84	2.3	2.51	1.04	2.73		
	HEM881002582	9.95	12.82	19.84	6.39	8.66	6.74	3.36	2.01	3.49	*	-
	HEM881002584	4.7	5.64	13.38	4.24	5.58	5.79	3.53	1.46	2.34		
45	HEM881002587	8.59	15.17	29.93	11.81	13.87	11.06	10.85	4.49	11.04		
	HEM881002590	27.91	31.34	78.12	45.05	22.1	24.36	21.48	22.81	39.4		
	HEM881002596	9.4	9.36	12.16	12.87	8.92	9.94	3.39	4.59	5.2	**	-
	HEM881002600	7.35	8.89	8.63	4.2	3.37	4.13	3.83	2.53	0.85	**	**
	HEM881002601	13.59	25.38	25.69	13.58	12.98	10.65	15.53	14.47	12.11		
	HEM881002603	4.01	4.35	8.49	3.09	3.23	2.33	3.4	1.13	1.07		
50	HEM881002607	5.59	8.75	13.05	6.25	8.3	3.46	5.04	2.83	4.29		
	HEM881002610	6.17	9.3	18.74	7.02	8.85	5.16	5.21	2.01	3.82		
	HEM881002613	5.1	6.85	16.57	4.47	6.62	3.43	4.07	1.86	3.63		
	HEM881002614	108.18	118.26	257.79	160.19	127.61	115.27	80.95	70.77	100.61		
	HEM881002615	3.76	4.74	6.46	3.53	4.89	3.11	2.31	1.58	0.85	*	-
	HEM881002617	15.34	16.43	20.69	14.06	10.46	7.36	16.03	9.38	12.23		
55	HEM881002623	43.46	62.43	70.11	89.32	59.95	64.13	46.57	40.8	51.16		
	HEM881002624	20.45	39.7	56.1	29.45	24.85	24.11	26.5	22.1	22.65		
	HEM881002631	12.8	24.72	22.26	5.67	13.71	7.87	12.49	14.8	9.64		

Table 388

5	HEM881001996	27.04	42.61	48.67	17.55	24.82	18.56	28.17	23.4	25.36	*	-
	HEM881001997	14.71	23.01	29.3	15.9	17.15	12.6	18.98	11.27	17.5		
	HEM881001999	15.2	22.46	20.51	13.68	17.9	14.36	19.76	8.32	11.63		
	HEM881002002	6.41	8.68	14.02	3.86	4.69	7.48	2.32	2.31	9.35		
	HEM881002005	13.93	14.9	24.54	14.07	12.64	13.23	24.2	15.6	24.4		
10	HEM881002009	13.81	20.37	28.5	12.9	13.24	9.2	14.63	11.53	16.62		
	HEM881002013	35.52	43.43	61.55	22.34	25.62	25.39	38.97	23.31	36.2	*	-
	HEM881002015	17.64	16.37	26.67	10.62	13.94	14.86	13.44	9	13.73		
	HEM881002024	5.8	10.6	13.79	2.3	5.04	3.13	2.77	1.85	3.6	*	-
	HEM881002035	52.65	85.52	32.75	21.71	70.63	38.58	40.44	20.7	17.84		
15	HEM881002039	4.95	5.73	12.99	4.99	10.04	5.87	7.91	2.61	3.57		
	HEM881002041	26.99	42.34	59.33	33.42	31.41	30.59	35.5	25.94	36.66		
	HEM881002042	7.32	8.11	15.44	8.43	5.37	4.7	6.95	4.45	6.64		
	HEM881002043	15.38	16.33	33.37	9.15	15.16	12.75	12.63	10.91	12.91		
	HEM881002044	6.28	7.19	15.99	4.63	6.03	3.49	8.95	3.61	5.78		
20	HEM881002045	13.55	14.05	38.81	21.11	17.68	16.3	14.27	7.83	17.46		
	HEM881002049	20.72	21.4	45.82	13.96	20.22	15.6	21.11	13.82	17		
	HEM881002050	12.83	16.25	37.04	11.52	18.46	11.6	18.41	9.54	14.5		
	HEM881002051	28.94	18.92	52.44	23.53	26.05	15.13	33.91	14.41	28.65		
	HEM881002068	64.56	61.13	134.22	53.82	39.3	49.08	48.07	48.15	30.52		
25	HEM881002069	20.88	27.08	36.85	26.67	17.79	18.95	23.6	16.68	23.73		
	HEM881002075	15.68	17.27	22.02	11.81	12.44	12.63	5.87	5.58	11.32	*	*
	HEM881002079	27.46	41.14	33.47	23.15	21.35	9.24	13.25	11.63	12.65	**	-
	HEM881002080	27.32	43.03	71.5	48.33	38.81	19.92	37.68	22.99	28.59		
	HEM881002082	11.54	12.93	24.86	5.24	21.72	15.36	15.59	4.71	14.1		
30	HEM881002084	17.19	20.49	48.44	15.57	24.56	30.61	24.16	14.34	22.35		
	HEM881002088	4.86	8.19	14.88	4.12	8	4.95	6.55	2.38	8.06		
	HEM881002092	144.34	116.11	135.03	45.77	56.49	60.46	15.5	25.51	16.94	**	**
	HEM881002094	15.12	19.26	35.12	17.76	15.07	8.88	14.96	9.23	22.31		
	HEM881002103	5.46	7.53	11.18	8.4	4.51	4.6	8.52	3.47	4.97		
35	HEM881002109	59.58	83.46	107.06	62.19	50.92	51.12	65.01	50.9	42		
	HEM881002115	5.46	60.72	14.77	39.13	16.84	5.85	5.92	6.74	5.3		
	HEM881002120	6.4	5.75	12.8	2.74	5.51	2.64	3.13	1.57	3.49		
	HEM881002121	19.45	25.43	55.52	28.56	24.08	19.83	25.58	15.89	20.17		
	HEM881002134	10.91	24.24	43.39	24.03	23.79	24.09	19.26	8.44	14.1		
40	HEM881002136	18.33	18.78	45.57	23.81	13.25	20.44	11.9	15.85	24.31		
	HEM881002138	50.09	51.59	110.11	35.51	42.15	39.79	27.69	28.58	25.22		
	HEM881002139	8.36	15.47	10.43	4.01	4.89	4.74	7.89	6.83	4.81	*	-
	HEM881002141	9.19	16.96	13.1	1.9	10.98	7	5.65	7.53	4.35	*	-
	HEM881002142	6.36	8.37	8.62	2.74	2.82	2.25	4.04	2.02	5.71	**	*
45	HEM881002145	10.49	16.24	32.26	9.14	12.19	10.34	12.54	5.18	7.82		
	HEM881002152	24.88	22.9	59.51	8.84	21.58	20.2	34.39	10.44	7.92		
	HEM881002162	46.61	80.75	107.44	66.24	81.5	44.12	82.89	47.69	70.76		
	HEM881002173	3.55	25.4	7.84	5.45	25.86	14.76	4.91	5.53	5.99		
	HEM881002189	5.91	5.36	11.91	4.49	4.74	3.73	7.69	4.5	2.15		
50	HEM881002190	8.05	5.19	7.81	3.82	3.51	3.24	3.4	1.73	2.03	*	*
	HEM881002193	13.5	31.42	40.95	12.51	15.71	29.81	30.17	23.34	31.46		
	HEM881002217	11.76	15.56	34.59	12.72	11.26	16.37	17.63	11.07	12.8		
	HEM881002218	25.74	52.63	73.43	32.82	28.59	24.52	35.6	20.64	25.88		
	HEM881002228	108.24	84.59	214.25	80.62	78.36	108.65	166.71	123.42	65.1		
55	HEM881002232	10.76	24.39	31.98	19.41	21.33	17.37	22.17	6.39	14.13		
	HEM881002245	29.32	39.47	46.03	33.69	25.42	30.62	28.25	20.36	26.6		
	HEM881002247	23.04	30.4	36.26	21.55	23.11	19.18	30.01	21.82	18.27		
	HEM881002249	8.45	5.4	12.9	9.1	8.18	7.58	2.92	2.96	4.18		
	HEM881002254	13.55	8.54	22.22	9.35	11.14	10.19	5.38	9.93	4.18		
55	HEM881002255	24	39.55	56.2	39.15	37.05	37.31	39.64	23.55	35.88		
	HEM881002266	19.55	37.2	69.12	27.4	30.03	37.09	27.79	17.42	23.42		
	HEM881002271	8.4	14.75	33.73	16.63	15.77	17.75	17.9	8.81	11.82		
	HEM881002280	8	1.32	3.2	1.59	2.57	0.19	2.26	0	1.09		
	HEM881002296	13.25	17.07	48.86	29.84	17.46	25.16	17	14.47	20.56		
55	HEM881002300	7.2	17.96	22.84	22.43	7.83	12.63	8.34	8.09	9.04		
	HEM881002302	8.19	9.95	13.55	11.87	6.29	9.78	7.21	2.24	3.19	*	-
	HEM881002306	4.46	4.94	10.04	8.42	4.47	9.84	5.18	3.92	6.24		
	HEM881002316	9.41	18.69	11	13.6	12.17	15.43	7.32	2.42	6.96		

Table 387

	HEM881001735	41.8	74.33	131.1	48.7	40.13	60.42	53.81	28.45	83.52
	HEM881001736	7.77	13.39	28.59	8.57	9.16	5.35	8.09	5.45	9.42
5	HEM881001747	2.69	8.82	13.53	4.41	6.88	3.98	4.96	1.05	3.48
	HEM881001749	10.28	24.28	26.11	5.15	8.2	10.2	5.99	4.27	10.01
	HEM881001753	10.3	11.05	20.9	9.25	9.8	9.74	6.82	5.36	10.12
	HEM881001756	6.53	8.08	12.86	3.9	3.99	4.72	5	3.97	3.65
	HEM881001757	7.8	24.09	26.61	6.4	12.83	5.96	11.18	14.36	9.41
10	HEM881001760	14.27	19.82	42.06	14.81	14.45	11.21	17.24	9.18	15.44
	HEM881001762	0.8	3.04	8.27	0.57	4.82	0.71	2.56	0.81	3.52
	HEM881001780	13.76	23.09	49.83	16.38	14.53	12.62	15.44	6.8	13.71
	HEM881001785	6.49	12.36	24.07	15.82	12.17	15.29	9.31	2.84	7.32
	HEM881001788	25.43	19.69	45.01	19	16.71	23.01	19.54	20.38	26.43
	HEM881001793	5.14	15.25	7.48	1.07	13.29	8	16.96	8.27	6.44
15	HEM881001797	14.78	27.69	37.34	12.45	18.06	22.63	22.37	14.75	11.18
	HEM881001802	13.95	6.42	12.11	6	8.1	3.62	4.89	7.62	4.73
	HEM881001812	19.94	26.82	51.97	30.77	23.99	24.59	25.74	15.27	19.71
	HEM881001815	11.81	15.53	37.64	17.99	17.45	14.83	20.39	13.54	19.43
	HEM881001816	17.77	23.19	75.37	26.22	24.8	18.06	30.17	15.34	21.15
20	HEM881001831	8.13	8.07	23.31	9.34	10.75	10.45	10.25	3.73	10.62
	HEM881001834	14.4	18	35.74	22.13	11.64	13.05	15.55	10.35	12.6
	HEM881001836	30.28	39.5	43.48	42.55	36.92	34.82	31.2	18.99	16.41
	HEM881001839	14.36	16.68	38.4	17.83	16.75	14.74	12.83	10.7	10.84
	HEM881001841	25.77	32.45	51.17	33.9	26.13	25.07	27.91	22.04	19.78
	HEM881001844	15.48	34.84	44.55	31	24.08	23.04	21.12	16.3	25.7
	HEM881001847	17.89	27.89	60.83	26.2	23.47	23.52	21.19	9.97	20.24
25	HEM881001848	4.85	5.43	11.29	3.58	4.51	3.65	6.07	3.08	5.61
	HEM881001850	38.14	52.12	78.88	56.18	68.55	53.58	61.43	34.94	43.46
	HEM881001859	30.81	27.05	59.36	68.19	35.46	57.28	19.8	14.26	20.24
	HEM881001863	3.46	3.98	8.32	5.64	4.36	4.8	7.82	2.45	4.74
	HEM881001867	14.5	16.62	28.09	12.79	10.98	12.92	14.13	8.77	15.8
	HEM881001868	5.84	8.18	12.72	7.12	6.24	5.61	8.09	8.94	8.48
30	HEM881001869	54.86	64.59	94.37	86.31	67.89	82.85	80.48	51.3	89.82
	HEM881001872	6.07	11.98	13.6	18.68	9.46	7.21	18.78	6.55	10.02
	HEM881001874	3	3.3	6.61	4.59	12.05	11.19	3.13	1.36	5.74
	HEM881001875	12.99	18.2	23.61	26.52	29.44	24.9	34.89	17.55	28.83
	HEM881001880	18.04	17.79	33.89	31.27	17.63	20.15	14.84	10.37	14.08
	HEM881001899	34.29	45.59	63.32	39.63	32.89	31.82	35.58	26.2	35.72
35	HEM881001903	8.16	10.41	29.09	8.47	10.89	12.17	9.48	4.66	7.91
	HEM881001905	28.16	29.19	50.5	33.46	32.78	32.37	33.69	21.9	33.71
	HEM881001906	4.65	5.05	9.2	6.27	6.88	4.65	7.25	6.26	9.65
	HEM881001908	2.42	5.7	6.4	5.21	4.63	6.7	4.19	0.96	5.33
	HEM881001910	5.2	10.62	10.84	8.51	10.51	9.98	6.79	2.76	7.39
40	HEM881001911	29.98	51.31	56.74	43.98	49.18	34.13	49.74	26.26	62.6
	HEM881001915	21.14	31.44	37.82	20.71	15.99	14.9	18.9	21.96	29.35
	HEM881001921	7.15	10.68	16.34	7.38	7.15	5.66	5.97	5.19	7.01
	HEM881001922	17.41	22.59	26.93	13.84	10.93	18.21	23.02	11.95	17.79
	HEM881001925	13.97	12.3	26.77	11.6	11.78	10.4	15.51	12.85	15.94
	HEM881001930	40.03	59.19	73.92	44.03	41.91	43.37	40.7	31.14	54.53
45	HEM881001944	1.94	5.58	8.4	6.66	4.5	5.16	2.67	2.16	4.4
	HEM881001945	11.9	14.15	28.25	12.61	15.86	30.85	15.45	10.36	15.75
	HEM881001947	12.74	2.15	16.64	14.03	14.01	12.25	15.71	4.99	12.24
	HEM881001950	17.57	25.59	40.32	25.51	11.85	16.5	13.53	12.65	19.84
	HEM881001952	26.14	33.12	47.22	23.32	21.94	24.22	34.78	37.99	49.1
	HEM881001953	17.2	26.64	33.2	20.18	17.54	17.43	18.02	10.46	24.03
50	HEM881001957	5.93	5.66	12.29	2.67	5.82	5.6	4.01	5.73	6.62
	HEM881001959	4.28	3.66	11.59	1.82	2.82	5.97	1.57	2.64	3.72
	HEM881001962	11.82	43.87	27.13	12.9	32.94	14.36	12.1	15.99	21
	HEM881001967	34.68	44.03	60.34	44.77	38.71	38.73	44.26	27.2	37.33
	HEM881001973	28.25	35.18	38.62	28.67	29.33	21.67	37.28	18.87	23.03
	HEM881001978	70.1	95.85	135.62	108.46	60.16	80.87	54.72	61.03	86.88
55	HEM881001983	4.69	4.06	8.04	3.84	5.22	5.54	5.15	3.31	5.57
	HEM881001987	10.55	12.27	28.1	12.58	12.32	11.95	11.85	6.54	23.57
	HEM881001988	9.33	7.22	23.4	8.24	11.34	8.88	13.31	6.58	9.69
	HEM881001990	26.2	19.68	27.76	43.3	26.68	36.79	27.71	13.13	24.3

Table 386

	HEM881001458	9.69	12.19	21.36	34.84	17.29	10.08	13.2	9.97	12.41			
	HEM881001461	19.67	23.1	39.56	20.5	28.42	24.75	17.54	14.4	25.15			
5	HEM881001463	19.56	21.33	35.61	31.93	33.77	43.28	30.72	20.68	36.42			
	HEM881001464	28.01	27.99	31	14.62	14.82	15.04	16.45	13.58	23.87	**	*	-
	HEM881001466	15.16	15.92	27.83	19.85	13.38	15.96	13.7	10.31	15.02			
	HEM881001482	17.23	20.81	35.32	19.01	19	19.25	19.83	14.79	21.88			
	HEM881001500	41.82	45.54	68.26	37.83	39.48	37.77	46.2	37.01	39.45			
10	HEM881001505	20.94	22.59	32.35	26.78	22.08	24.15	26.64	17.19	32.59			
	HEM881001521	8.53	7.82	10.6	8.16	8.3	6.64	6.41	3.73	7.14			
	HEM881001527	2.99	3.88	4.38	4.6	7.15	3.37	3	1.09	4.09			
	HEM881001530	3.48	3.01	4.27	1.8	4.75	3.13	7.05	4.01	3.15			
	HEM881001531	36.17	50.68	53.86	24.43	15.05	19.88	14.37	19.2	25.31	*	*	-
	HEM881001532	17.53	24.2	32.45	18.39	15.75	13.6	15.22	13.76	17.31			
15	HEM881001535	12.38	16.91	23.89	12.47	13.36	10.82	9.04	12.27	13.78			
	HEM881001536	66.38	84.8	143.72	91.2	78.03	69.03	101.64	71.92	118.47			
	HEM881001537	11.79	11.39	16.93	9.29	11.02	10.91	6.73	5.87	10.85			
	HEM881001542	14.96	22.23	29.71	13.95	16.64	16.35	12.71	5.92	15.48			
	HEM881001543	23.66	28.65	30.69	20.28	24.21	31.58	17.34	19.58	58.07			
	HEM881001547	23.94	50.04	39.45	14.74	15.27	12.72	12.08	2.84	4.52	*	*	-
20	HEM881001548	3.11	4.4	5.6	1.9	1.94	2.29	3.65	4.19	3.74	*	*	-
	HEM881001551	3.8	7.07	10.09	4.64	4.24	4.26	3.14	1.65	6.16			
	HEM881001555	8.19	9.59	15.51	4.72	6.09	5.64	3.45	5.22	5.59			
	HEM881001562	17.05	18.48	38.18	19.29	13.39	17.57	16.22	12.15	22.52			
	HEM881001564	38.37	50	64.37	49.41	38.27	48.76	40.56	25.46	46.57			
	HEM881001565	18.11	21.08	35.2	23.9	18.29	10.69	18.09	9.6	15.5			
25	HEM881001569	18.83	22.86	28.98	20.6	18.61	17.83	18.18	11.82	14.18			
	HEM881001573	3.68	5.99	8.15	7.26	5.87	2.52	10.81	4.11	5.2			
	HEM881001585	31.55	42.11	57.35	32.5	28.22	32.05	31.45	26.22	33.57			
	HEM881001586	19.62	27.08	60.2	25.12	17.34	21.21	21.42	13.56	26.31			
	HEM881001588	3.6	1.4	8.8	2.53	0.56	1.81	2.28	0.36	3.78			
	HEM881001595	10.36	11.37	25.05	8.69	10.88	9.92	9.72	7.65	13.65			
30	HEM881001596	50.23	70.48	94.67	85.43	78.99	83.08	60.67	47.9	57.78			
	HEM881001598	12.08	19.55	22.95	3.76	11.11	7.77	4.14	3.06	5.8	*	*	-
	HEM881001603	15.69	23.5	36.56	18.96	19.27	14.23	19.02	14.12	17.69			
	HEM881001606	8.92	10.53	14.14	3.61	11.6	8.26	6.12	2.83	3.66	*	*	-
	HEM881001612	12.97	18.3	27.94	17.56	18.03	14.45	13.69	13.84	21.14			
	HEM881001618	47.24	51.05	53.93	31.05	43.7	40.99	38.7	45.72	63.89	*	*	-
35	HEM881001619	21.53	28.8	35.45	21.69	15.6	18.53	19.99	9.27	17.16			
	HEM881001623	32.27	35.63	43.91	23.19	18.98	32.29	44.61	25.32	56.22			
	HEM881001625	8.57	7.58	12.04	5.34	5.5	6.91	4.26	3.63	6.59	*	*	-
	HEM881001630	10.5	14.13	23.29	7.12	9.43	7.75	11.72	5.74	8.87			
	HEM881001635	8.75	10.43	14.63	5.84	10.39	7.88	13.65	6.82	10.88			
	HEM881001637	32.58	33.36	51.7	33.85	36.88	33.36	45.82	19.62	25.12			
40	HEM881001641	50.01	60.79	92.28	48.36	49.98	43.85	47.93	38.87	56.74			
	HEM881001653	12.57	22.18	30.48	17.59	15.29	13.82	24.06	13.06	30.12			
	HEM881001665	3.93	2.53	9.42	3.83	1.95	2.69	2.78	1.9	1			
	HEM881001666	19.92	21.89	36.37	13.51	17.12	14.78	11.97	10.86	17.21			
	HEM881001667	16.89	20.47	51.29	17.31	18.04	14.47	10.43	9.14	14.45			
	HEM881001668	4.06	3.82	13.97	2.65	3.74	1.89	2.27	0.07	2.33			
45	HEM881001689	4.2	4.89	17.86	5.72	8.1	3.19	6.26	1.74	4.46			
	HEM881001670	46.37	46.73	48.84	13.56	78.84	42.13	34.32	15.02	19.72	*	*	-
	HEM881001673	67.96	61.75	141.78	71.26	44.3	44.21	62.82	56.55	66.02			
	HEM881001675	17.04	14.33	30.97	18.1	10.45	10.58	21.47	8.1	12.88			
	HEM881001679	6.24	17.91	21.96	5.19	4.96	3.87	10.86	6.26	8.43			
	HEM881001684	10.61	21.58	17.28	4.97	9.8	7.91	15.07	5.39	2.51			
50	HEM881001685	5.57	8.99	14.97	3.84	6.75	3.87	5.52	1.85	2.86			
	HEM881001695	5.09	9.94	17.45	5.82	8.3	5.19	6.66	2.16	2.97			
	HEM881001703	4.68	6.84	16.43	5.87	6.69	4.05	8.72	2.46	4.64			
	HEM881001704	5.61	7.88	21.01	10.94	17.96	9.05	14.4	6.3	9.45			
	HEM881001706	7.42	2.91	5.37	4.07	5.66	4.18	6.33	3.36	3.19			
	HEM881001707	3.87	5.86	5.44	3.17	4.08	5.01	5.27	1.48	7.88			
55	HEM881001717	14.88	12.76	30.35	13.57	11.78	8.58	13.5	11.39	12.48			
	HEM881001731	39.25	68.03	95.93	39.11	36.32	32.38	46.55	35.06	44.86			
	HEM881001734	5.17	6.48	12.05	1.52	7.19	2.49	5.94	1.69	6.44			

Table 385

	HEMB81001200	13.2	15.34	26.9	9.43	12.58	11.64	13.55	8.01	17.32			
	HEMB81001208	18.63	26.31	21.34	16.48	23.59	8.33	19.57	7.02	6.33			
5	HEMB81001209	6.61	9.43	16.76	4.47	13.17	7.28	9.02	4.73	7.13			
	HEMB81001210	2.22	5.57	9.4	5.43	2.97	2.71	2.73	1.63	4.83			
	HEMB81001215	9.7	12.03	15.37	9.6	10.02	9.52	8.36	4.91	10.29			
	HEMB81001217	10.51	10.25	17.62	6.77	6.49	7.62	10.25	4.74	16.86			
	HEMB81001218	6	5.57	10.2	5.12	3.8	3.11	4.86	3.23	6.96			
10	HEMB81001221	26.87	26.88	47.38	27.34	25.86	18.24	28	17.58	20.46			
	HEMB81001224	16.71	16.42	17.84	3.84	6.91	9.28	7.94	2.93	4.93	**	**	-
	HEMB81001230	17.23	21.95	42.64	15.78	20.28	10.72	28.16	18.31	25.4			
	HEMB81001234	6.94	12.6	13.24	4.58	8.31	8.14	5.53	1.54	8.42			
	HEMB81001235	4.87	3.91	1.57	0	1.12	0.17	1.32	0.88	4.29	*		-
	HEMB81001237	5.75	2.45	7.86	3.45	3.48	1.87	2.9	1.6	4.06			
15	HEMB81001242	16.88	13.35	16.55	9.1	8.16	9.95	11.54	6.31	10.57	**	*	-
	HEMB81001244	35.65	49.79	60.5	32.41	25.43	27.28	45.76	28.88	37.22			
	HEMB81001249	22.08	31.03	56.18	21.17	26.84	20.21	27.38	14.91	24.06			
	HEMB81001253	28.9	34.49	70.42	28.32	28.95	21.57	41.19	24.41	34.07			
	HEMB81001254	7.43	6.21	17.96	4.5	6.62	3.53	4.22	0.83	3.95			
20	HEMB81001266	69.19	46.43	92.09	93.11	71.97	69.72	54.64	34.46	57			
	HEMB81001267	104.19	98.07	152.36	13.92	33.06	57.1	34.39	45.85	70.55	*	*	-
	HEMB81001271	34.55	33.29	51.36	34.34	20.46	23.71	30.23	24.87	29.48			
	HEMB81001282	9.97	10.71	15.91	7.88	6.97	9.33	8.95	7.26	10.69			
	HEMB81001287	3.47	3.55	8.03	3.1	1.84	1.48	2.02	1.47	1.7			
	HEMB81001288	15.95	79.12	35.23	25.95	31.84	28.36	30.82	15.48	12.44			
	HEMB81001289	22.37	27.51	58.3	21.33	26.2	17.4	27.36	15.96	26			
25	HEMB81001290	4.47	6.46	15.08	5.02	4.85	3	3.41	2.52	3.43			
	HEMB81001294	300.12	442.69	478.77	294.44	431.41	291.22	536.94	211.14	291.4			
	HEMB81001299	11.76	27.27	23.84	14.94	18.67	41.4	14.69	12.85	20.6			
	HEMB81001302	14.9	20.75	33.76	16.74	10.48	13.15	15.46	13.18	16.41			
	HEMB81001304	20.54	137.19	27.35	22.15	29.27	12.7	29.77	14.5	13.73			
	HEMB81001314	6.45	11.88	16.63	9.21	7.91	6.84	9.36	5.15	7.34			
30	HEMB81001315	4.17	4.2	11.99	3.59	3.25	2.94	5.12	2.66	3.24			
	HEMB81001317	58.08	165.69	115.47	26.35	66.22	33.54	41.88	11.39	13.42	*		-
	HEMB81001326	7.12	11.9	25.13	9.45	10.76	5.58	7.57	2.53	8.72			
	HEMB81001331	60.43	105.03	131	66.61	74.76	59.35	124.09	50.05	41.99			
	HEMB81001335	0.53	0.14	1.28	1.92	0.74	0.41	1.38	0	1.24			
	HEMB81001337	19.31	21.8	38.53	21.67	17.28	16.88	20.3	14.12	26.23			
35	HEMB81001339	6.1	5.5	14.98	6.52	5.57	3.43	4.24	2.59	4.91			
	HEMB81001344	11.08	7.69	21.58	3.98	10.13	4.45	6.01	7.37	4.42			
	HEMB81001346	14.51	8.09	19.68	7.72	9.53	5.04	11.31	3.15	5.25			
	HEMB81001348	2.18	3.12	12.35	0.43	2.84	1.52	3.82	0.54	2.87			
	HEMB81001350	52.05	108.41	129.66	78.51	93.58	81.19	64.58	43.45	65.59			
40	HEMB81001356	7.52	9.31	26.43	13.44	18.12	16.4	11.29	5.29	11.49			
	HEMB81001364	8.27	9.27	21.55	13.04	8.88	12.02	9.82	4.71	9.74			
	HEMB81001366	11.82	16.33	29.77	17.62	9.11	9.15	17.69	10.49	12.45			
	HEMB81001367	34.9	33.46	79.86	33.05	29.6	26.79	45.62	23.04	29.37			
	HEMB81001369	13.59	34.47	15.16	9.61	17.82	9.58	10.33	11.36	4.3			
	HEMB81001380	5.78	6.69	14.57	4.06	3.91	3.5	3.91	2.87	4.28			
	HEMB81001381	7.8	18.4	21.36	7.96	16.58	12.19	13.62	2.72	8.14			
45	HEMB81001384	64.99	90.97	10.61	0.27	3.72	0.15	67.41	60.83	3.42			
	HEMB81001387	12.67	18.5	30.56	17.22	19.87	15.36	19.61	9.85	12.99			
	HEMB81001394	1.26	2.69	4.65	6.43	2.24	3.63	0.47	1.11	0.58			
	HEMB81001407	7.9	11.42	30.12	18.8	14.97	11.43	10.87	6.27	10.35			
	HEMB81001410	2.04	3.94	3.61	2.8	2.74	2.85	1.42	0.21	2.14			
	HEMB81001413	4.48	4.13	8.85	5.53	3.83	5.6	1.34	2.67	1.9			
50	HEMB81001419	11.47	23.84	24.98	19.26	14.85	15.61	11.95	7.31	16.96			
	HEMB81001421	5.76	4.41	16.42	7.89	11.28	6.94	5.16	2.14	5.02			
	HEMB81001424	2.91	4.4	4.11	0.76	0	0.56	5.88	1.84	0.07	**		-
	HEMB81001426	14.06	23.97	51.32	27.45	25.66	29	18.89	8.5	8.91			
	HEMB81001429	6.88	11.32	20.01	14.76	14.71	20	10.35	6.2	12.32			
	HEMB81001436	18.71	17.94	33.05	24.24	11.69	16.55	21.47	12.89	21.69			
55	HEMB81001443	19.58	21.95	43.4	20.44	16.81	19.91	22.72	13.56	24.01			
	HEMB81001449	35.58	35.95	71.7	35.81	35.7	39.37	42.62	30.92	51.15			
	HEMB81001454	4.44	10.09	10.48	10.32	5.73	8.72	4.72	2.95	5.93			

Table 384

	HEMB81000915	26.06	60.99	57.47	25.29	36.05	22.49	27.81	17.27	27.94			
5	HEMB81000917	5.69	5.32	12.64	3.03	4.44	3.24	2.96	0.58	2.57			
	HEMB81000927	17.9	20.69	49.41	20.99	18.29	12.76	18.76	15.57	19.4			
	HEMB81000932	14.04	31.9	58.79	23.84	30.24	22.56	24.51	13.79	19.35			
	HEMB81000933	73.2	82.97	106.78	17.69	17.17	51.59	30.19	46.68	77.12	*	-	
	HEMB81000936	28.64	37.39	39.34	11.49	17.42	18.08	9.46	19.76	21.12	**	*	-
	HEMB81000939	13.01	16.47	24.62	8.79	10.18	11.8	13.33	5.75	13.18			
10	HEMB81000941	55.55	67.95	96.78	34.29	51.94	34.02	64.38	50.22	47.74			
	HEMB81000947	13.1	15.93	33.36	11.69	15.06	7.82	14.81	8.66	11.7			
	HEMB81000954	9.7	14.84	29.47	10.7	10.34	11.24	13.58	8.18	14.09			
	HEMB81000959	5.31	7.05	16.81	1.63	2.59	1.53	2.83	0	2.89			
	HEMB81000973	7.09	9.47	31.83	11.47	11.07	10.69	11.63	4.69	9.55			
	HEMB81000975	3.47	3.38	0.39	0.25	0.11	0.78	2.64	0.48	3.5			
15	HEMB81000981	13.64	11.74	28.6	8.77	7.91	9.41	10.88	6.85	13.87			
	HEMB81000985	21.62	19.08	51.64	22.07	25.82	18.12	29.3	14.45	17.75			
	HEMB81000991	46.57	60.97	104.38	49.12	52.31	37.89	58.02	46.82	42.32			
	HEMB81000996	7.55	10.13	19.44	15.43	7.58	7.76	25.49	7.93	12.48			
	HEMB81001000	8.66	10.27	29.46	13.94	12.61	10.4	13.32	7.07	9.02			
	HEMB81001004	4.52	20.53	22.05	7.15	7.85	4.71	7.51	4.91	9.37			
20	HEMB81001008	27.45	57.49	106.29	36.82	29.09	41.72	54.86	25.88	43.51			
	HEMB81001011	13.92	18.59	22.13	8.85	6.12	13.6	9.78	8.17	13.65			
	HEMB81001014	4.93	5.88	7.48	5.94	4.03	6.17	3.66	1.76	2.96	*	-	
	HEMB81001020	38.49	45.64	94.13	52.12	43.16	44.7	42.48	28.95	25.72			
	HEMB81001024	14.35	17.88	33.12	21.84	15.86	15.33	12.31	9.54	11.76			
	HEMB81001026	36.33	37.82	64.87	35.84	31.03	31.81	34.99	27.91	28.23			
25	HEMB81001037	29.37	38.91	99.73	42.44	38.61	41.96	49.51	23.14	48.42			
	HEMB81001042	40.43	19.68	79.17	12.87	30.43	38.57	43.54	27.06	24.73			
	HEMB81001046	1.02	3.17	4.37	2.13	2.63	1.33	3.19	0	2.81			
	HEMB81001047	3.32	3.99	7.05	6.87	5.14	9.57	4.92	2.39	5.02			
	HEMB81001048	14.93	18.87	36.47	38.03	15.11	21.97	9.91	8.51	13.37			
	HEMB81001051	16.67	10.51	26.31	24.66	18.42	23.37	16.85	8.02	15.72			
30	HEMB81001056	5.76	6.07	15.19	9.62	10.04	7.15	7.61	2.99	9.72			
	HEMB81001058	13.02	13.37	20.82	14.04	12.97	13.5	13.64	8.33	17.02			
	HEMB81001060	6.48	6.79	11.56	3.09	7.6	8.52	5.42	2.73	5.08			
	HEMB81001063	7.02	9.34	7.27	6.14	6.79	5.01	6.52	5.39	11.16			
	HEMB81001068	12.88	13.51	18.25	11.13	9.67	11.92	8.97	9.12	12.67			
	HEMB81001082	19.71	21.28	25.07	26.2	26.41	15.74	14.59	12.53	27.85			
35	HEMB81001095	9.14	8.65	18.5	8.62	9.69	8.93	5.69	6.23	9.72			
	HEMB81001096	44.08	45.29	88.46	46.23	37.21	42.49	31	20.55	25.13			
	HEMB81001101	14.24	13.37	19.97	9.53	14.04	14.7	13.73	9.26	15.99			
	HEMB81001102	5.29	6.01	11.42	4.54	5.69	5.17	5.82	3.61	6.55			
	HEMB81001104	38.08	39.86	49.87	41.44	36.7	31.13	23.82	16.12	29.93	*	-	
	HEMB81001105	44.24	54.5	42.57	26.76	34.74	25.96	40.78	32.97	47.69	*	-	
40	HEMB81001112	7.32	12.27	12.46	6.67	5.72	6.06	2.52	6.54	5.01	*	-	
	HEMB81001113	12.36	16.73	27.57	10.87	11.17	11.23	11.03	6.35	13.81			
	HEMB81001114	10.5	25.04	16.53	8.31	9.62	7.81	7.26	6.53	13.76			
	HEMB81001115	5.38	5.54	12.69	5.57	5.33	6.84	5.79	8.01	10.93			
	HEMB81001117	24	32.48	48.34	30.8	26.94	25.33	34	33.8	46.28			
	HEMB81001119	6.59	8.79	10.75	5.08	5.94	5.54	5.24	3.59	5.78	*	-	
45	HEMB81001126	22.85	31.95	38.8	19.87	19.28	19.37	24.09	9.52	26.55			
	HEMB81001133	2.49	4.42	8.71	2.43	5.35	3.57	4.22	0.99	2.59			
	HEMB81001137	7.73	11.05	0	0.73	0.66	0.68	7.84	6.26	16.01			
	HEMB81001142	3.94	3.85	6.48	2.73	4.14	2.88	2.5	3.32	8.04			
	HEMB81001145	7.24	10.68	17.23	5.39	7.36	5.26	4.34	3.98	7.6			
	HEMB81001151	3.26	6.63	6.67	3.5	3.71	2.34	0.95	2.94	4.05			
50	HEMB81001153	3.1	3.77	9.68	1.47	2.32	1.7	2.21	0	1.53			
	HEMB81001158	5.14	4.68	9.64	3.27	3.84	3.55	2.97	1.38	2.99			
	HEMB81001169	31.33	46.44	55.42	27.9	34.88	25.03	41.78	26.83	37.34			
	HEMB81001170	8.62	15.69	19.05	7.59	11.25	8.63	12.14	5.58	9.38			
	HEMB81001175	2.65	3.14	5.1	4.52	2.61	4.91	2.72	1.86	6.12			
	HEMB81001177	5.44	6.28	5.09	1.64	3.43	6.94	2.77	2.32	6.91			
55	HEMB81001182	18.97	18.18	22.47	9.28	11.35	10.35	7.83	8.39	8.5	**	**	-
	HEMB81001192	9.02	9.64	12.65	5.82	4.73	5.18	6.79	2.78	10.91	*	-	
	HEMB81001199	4.71	4.4	13.79	2.26	2.76	5.59	3.91	3.02	2.96			

Table 383

	MEMB81000631	17.85	20.22	28.17	18.29	17.71	18	15.1	13.03	22.53			
	MEMB81000632	9.09	8	17.51	9.07	9.79	9.7	5.89	4.13	8.31			
5	MEMB81000636	2.41	3.55	6.37	2.23	5.22	2.73	5.28	1.01	4.17			
	MEMB81000637	10.09	17.58	15.59	10.48	18.82	18.4	10.07	3.67	7.99			
	MEMB81000638	2.99	4.17	4.52	3.32	2.17	3.48	2.49	1.75	2.68	*		
	MEMB81000642	39.5	47.36	81.46	47.61	39.34	45.01	39.82	28.49	56.57			
	MEMB81000643	19.66	15.32	32.08	23.95	12.4	14.81	11.1	10.59	13.67			
	MEMB81000649	12.99	15.1	28.14	16	12.05	14.32	13.77	8.35	17			
10	MEMB81000652	14.35	8.01	17.67	7.91	13.41	10.82	9.88	9.25	11.18			
	MEMB81000655	4.61	7.58	7.57	5.6	5.2	3.31	4.16	0.72	5.12			
	MEMB81000665	4.16	6.24	9.15	3.47	7.38	6.78	5.21	1.62	5.11			
	MEMB81000668	16.93	27.34	0.25	1.82	1.14	0.36	23.87	15.76	19.86			
	MEMB81000671	38.43	46.48	92.78	44.14	31.73	40.3	25.53	28.64	41.28			
	MEMB81000673	32.99	36.97	43.71	17.4	18	19.87	6.02	26.42	8.4	**	*	-
15	MEMB81000679	47.01	69.39	88.27	46.41	43.74	44.85	52.81	42.78	53.45			
	MEMB81000684	6.8	10.16	24.24	9.32	9.77	12.54	6.96	8.48	13.11			
	MEMB81000692	16.03	14.55	25.59	15.28	15.92	17.55	16.09	8.5	14.57			
	MEMB81000693	1.76	2.64	8.52	1.53	4.27	2.95	4.06	0.19	6.13			
	MEMB81000705	14.59	17.08	31.29	17.89	15.24	18.52	13.25	7.08	17.6			
20	MEMB81000706	11.29	12.22	20.56	12.35	17.3	12.09	6.4	4.7	7.15	*		-
	MEMB81000709	45.75	62.83	118.05	70.74	40.37	58.09	48.11	45.53	67.69			
	MEMB81000714	31.43	34.85	47.38	12.83	21.37	16.81	5.14	8.31	8.35	*	**	-
	MEMB81000725	5.12	8.87	10.38	4.23	3.1	4.36	4.23	1.66	4.96			
	MEMB81000726	0.86	1.95	5.91	0.06	0.38	0.65	0.77	3.05	1.2			
	MEMB81000729	6.42	9.45	14.46	7.38	5.18	5.16	5.32	4.5	6.56			
25	MEMB81000738	18.18	14.39	28.4	16.66	7.78	8.45	31.84	5.87	7.73			
	MEMB81000749	19.05	25.67	36.38	23.07	19.8	20.09	17.18	15.06	19.35			
	MEMB81000763	2.4	3.95	6.27	1.43	7.01	5.93	4.26	0.57	3.97			
	MEMB81000770	32.56	40.83	87.59	46.89	28.6	43.89	33.53	25.96	52.34			
	MEMB81000774	35.04	53.35	80.31	43.91	38.96	45.26	39.09	37.25	56.98			
	MEMB81000777	45.62	74.38	70.43	17.25	25.76	25.38	18.75	21.39	27.43	*	*	-
30	MEMB81000781	6.33	5.26	11.47	6.25	5.6	4	3.41	2.88	6.03			
	MEMB81000788	18.37	23.49	32.23	14.46	15.9	16.37	22.44	12.27	19.59			
	MEMB81000789	23.79	24.39	34.32	14.31	19.85	15.3	18.54	10.27	14.08	*	*	-
	MEMB81000790	5.59	8.56	17.48	2.6	6	2.29	4.26	1.79	6.06			
	MEMB81000794	18.79	29.68	40.8	14.17	25.07	14.74	20.31	17.36	24.13			
	MEMB81000807	12.94	8.25	2.03	1.02	2.13	2.36	2.69	4.28	6.87			
35	MEMB81000809	78.77	111.21	120.34	77.13	62.86	41.38	80.78	69.07	111.98			
	MEMB81000810	10.52	9.56	13.14	4.85	2.89	5.55	9.37	2.32	10.71	**		-
	MEMB81000821	11.49	9.52	21.32	7.91	10.41	6.87	11.4	5.99	7.83			
	MEMB81000822	14.47	17.81	31.19	12.64	15.48	12.05	15.29	6.81	15.5			
	MEMB81000826	7.31	8.22	12.01	4.01	6.22	5.16	2.28	1.16	3.63	*		-
	MEMB81000827	3.32	10.97	13.01	2.13	2.92	3.29	2.14	1.1	3.35			
40	MEMB81000831	2664.2	3968.9	2993.4	2288.3	2955.9	3122.5	3699.5	1769.3	603.46			
	MEMB81000835	65.9	87.92	134.3	99.52	82.05	71.78	46.56	49.76	78.94			
	MEMB81000840	22.85	22.57	43.26	24.08	23.34	18.79	20.99	18.48	32.32			
	MEMB81000848	11.78	12.27	18.41	9.27	11.15	6.47	11.75	6.28	10.74			
	MEMB81000852	6.96	7.13	13.91	5.13	5.29	6.06	3.72	3.05	2			
	MEMB81000857	14.9	19.2	33.57	16.67	15.78	14.82	14.34	9.21	17.14			
45	MEMB81000858	18.62	21.62	40.52	17.01	24.14	29.4	19.12	14.81	25.58			
	MEMB81000867	3.35	5.16	11.87	1.01	3.43	0.49	5.8	0.59	7.29			
	MEMB81000870	5.48	10.65	17.84	3.33	10.48	4.74	3.16	0.41	3.45			
	MEMB81000876	9.67	10.87	29.49	19.7	15.97	14.54	9.25	6.54	15.24			
	MEMB81000881	16.87	17.18	33.12	17.09	13.5	10.07	12.12	11.55	18.55			
	MEMB81000883	6.22	6.79	8.96	2.61	2.5	1.97	4.54	1.83	3.25	**	*	-
50	MEMB81000887	5.96	5.57	6.2	3.69	2.17	1.88	2.97	1.27	2.62	**	**	-
	MEMB81000888	3.12	5.25	10.45	4.39	3.56	2.35	3.24	2.38	1.97			
	MEMB81000890	10.06	35.98	23.82	9.78	19.94	41.91	13.4	7.59	9.49			
	MEMB81000893	34.65	25.07	51.31	20.43	19.67	19.95	21.33	13.24	12.88			
	MEMB81000900	14.11	15.32	20.53	11.04	14.54	9.38	11.56	4.15	3.3	*		-
	MEMB81000905	8.7	11.8	4.84	3.81	3.66	3.74	10.34	4.58	8.59			
55	MEMB81000908	10.34	11.07	11.16	10.97	10.92	24.1	19.82	7.35	8.29			
	MEMB81000910	14.49	13.52	29.52	12.54	13.67	12.24	17.07	12.01	16.77			
	MEMB81000913	52.28	41.72	87.34	40.16	51.04	32.05	56.11	40.33	43.91			

Table 382

	HEM881000332	4.52	4.05	10.22	2.71	3.85	3.38	4.09	2.98	4.92			
	HEM881000335	4.77	4.71	13.27	5.2	6.34	6.3	5.06	1.38	4.17			
5	HEM881000336	4.61	9.32	11.37	1.79	3.32	1.88	2.23	1.53	1.7	*	*	-
	HEM881000337	29.33	35.43	67.58	48.11	46.55	55.37	21.92	22.71	26.93			
	HEM881000338	34.7	54.08	50.89	33.23	39.29	35.27	51.12	24.63	33.57			
	HEM881000339	10.36	11.59	21.89	13.7	11	9.9	8.4	6.43	14.47			
	HEM881000341	7.15	6.94	11.5	5.95	5.91	6.13	8.18	5.44	9.71			
10	HEM881000343	22.54	22.93	36.07	24.96	20.99	25.34	24.47	20.51	25.26			
	HEM881000354	24.93	29.93	37.49	16.95	17.28	19.25	24.74	17.57	19.26	*		-
	HEM881000358	4.2	3.56	9.08	2.24	3.75	2.51	2.41	0	3.07			
	HEM881000369	7.77	9.59	8.26	0	3.16	0.74	4.97	3.09	8.59	**		-
	HEM881000373	4.9	5.6	9.86	3.73	5.67	3.02	3.88	2	4.12			
	HEM881000374	33.91	39.77	79.58	37.61	43.98	33.27	46.4	29.37	24.06			
15	HEM881000376	23.61	37.46	51.78	28.25	29.92	24.65	24.75	22.13	31.75			
	HEM881000383	18.87	23.34	28.75	7.22	8.14	5.48	8.08	3.98	4.53	**	**	-
	HEM881000391	18.94	23.82	39.78	18.85	17.85	18.64	26.87	13.66	21.63			
	HEM881000399	7.91	9.01	13.97	2.99	3.3	3.33	3.37	1.8	1.69	*	*	-
	HEM881000402	5.66	7.15	13.91	3.49	5.23	2.79	6.8	2.16	2.32			
	HEM881000404	4.49	5.13	14.15	2.37	3.5	4.1	3.34	1.25	1.09			
20	HEM881000407	10.22	16.01	27.45	10.21	12.59	7.8	5.46	2.69	5.89			
	HEM881000420	8.96	13.82	23.72	10.31	11.56	10.07	12.86	5.36	9.15			
	HEM881000430	87.5	75.56	120.02	34.45	32.4	42.69	83.02	74.93	33	*		-
	HEM881000434	34.1	37.7	48.77	30.73	22.51	21.34	32.9	22.77	33.21	*		-
	HEM881000438	4.47	5.36	6.73	3.08	3.98	3.03	2.33	1.63	2.04	*	**	-
	HEM881000441	14.04	26.42	32.37	16.04	15.64	13.8	26.09	16.54	18.05			
25	HEM881000447	26.11	49.46	42.13	39.56	42.28	10.3	26.83	15.38	20.58			
	HEM881000449	4.61	5.17	16.11	3.66	5.27	3.1	5.36	1.18	6.41			
	HEM881000453	36.46	39.41	48.95	17.29	27.48	11.55	15.42	4.83	8.75	*	**	-
	HEM881000455	7.98	15.4	31.12	12.14	13.6	12.23	19.02	4.77	15.17			
	HEM881000472	8.9	8.65	20.38	10.03	8.41	7.69	7.24	3.85	5.6			
	HEM881000480	18.47	20.31	37.47	19.05	17.04	11.84	20.16	15.95	20.84			
30	HEM881000486	36.69	36.48	47.19	21.3	19.24	16.88	39.82	23.52	22.96	**		-
	HEM881000487	6.37	6.26	12.27	7.11	4.47	4.08	6.03	4.45	5.18			
	HEM881000490	32.85	71.05	75.4	38.48	33.85	25.5	47.17	24.97	28.6			
	HEM881000491	10.98	13.88	21.94	11.39	11.14	7.83	9.44	3.7	10.75			
	HEM881000492	11.93	15.38	18.28	8.6	7.89	5.65	7.79	2.28	4.01	*	*	-
	HEM881000493	3.37	12.88	19.63	10.93	27.47	71.09	13.51	1.84	8.84			
35	HEM881000510	11.39	18.63	31.92	20.35	17	14.3	15.03	12.6	21.32			
	HEM881000516	5	6.48	9.48	3.37	2.52	5.44	4.17	3.02	4.74			
	HEM881000518	5.79	7.56	16.99	3.63	4.25	4.51	8.74	3.95	5.88			
	HEM881000523	16.51	22.61	33.14	14.54	14.09	13.74	17.51	11.52	13.77			
	HEM881000530	6.45	6.62	14.38	4.57	5.23	3.03	6.95	3.14	3			
40	HEM881000542	11.01	10.05	25.81	5.58	5.42	6.88	5.64	1.26	2.47			
	HEM881000550	7.5	9.67	18.88	5.5	6.7	4.85	3.66	1.58	2.78			
	HEM881000554	23.66	38.54	60.71	31.87	34.21	23.57	36.99	19.84	22.96			
	HEM881000556	5.71	9.66	13.39	4.65	5.23	10.47	3.94	4.86	6.74			
	HEM881000564	7.31	9.82	18.45	11.61	10.28	10.62	15.74	5.76	7.4			
	HEM881000567	20.61	20.17	47.53	8.75	7.89	6.38	13.88	8.67	4.74			
	HEM881000569	5.63	6.5	12.71	2.7	3.07	4.45	4.1	3.02	2.02			
45	HEM881000573	32.51	36.2	72.29	43.4	30.21	37.17	37	20.09	25.58			
	HEM881000575	14.84	26.39	42.18	17.14	17.78	15.29	22.14	11.44	16.53			
	HEM881000579	12.25	14.06	28.61	6.45	6.01	11.92	12.64	5.71	5.94			
	HEM881000585	4.75	6.58	10.45	5.85	6.89	6.63	6.35	0.91	4.03			
	HEM881000586	9.22	12.19	32.5	16.92	12.87	17.45	7.51	6.24	9.01			
	HEM881000589	11.66	16.54	23.61	18.44	15.28	16.1	11.67	7.91	7.82			
50	HEM881000591	12.71	17.74	34.25	24.96	19.88	19.6	12.6	9.24	10.66			
	HEM881000592	4.57	3.51	5.52	1.66	2.6	2.47	2.07	3.49	1.32	*		-
	HEM881000593	91.28	145.08	255.41	284.78	269.98	262.29	122.51	62.28	150.94			
	HEM881000595	28.02	34.64	53.56	11.28	18.32	41.15	22.88	29.26	17.38			
	HEM881000598	22.01	31.15	65.04	24.88	33.54	29.35	68.32	20.78	23.46			
	HEM881000611	2.41	5.2	8.28	7.8	5.78	3.85	5.15	1.83	3.38			
55	HEM881000617	53.65	47.37	106.59	110.75	55.04	84.65	42.52	37.92	47.84			
	HEM881000623	31.42	31.62	79.25	50.87	28.93	40.28	29.97	27.06	51.77			
	HEM881000630	5.75	17.04	13.07	8.82	8.82	8.76	7.96	4.83	11.56			

Table 381

	HEMBA1007347	16.98	22.65	29.56	14.57	13.83	11.3	16.82	11.86	19.8			
	HEMBA1007353	5.79	10.55	11.68	3.83	9.63	8.03	10.26	4.85	8.24			
5	HEMBA1000095	12.22	17.75	30.52	11.82	12.78	11.43	13.73	6.39	17.44			
	HEMBA1000098	18.71	35.13	41.01	19.83	22.41	13.75	23.85	15.07	25.11			
	HEMBA1000018	26.09	49.37	73.92	38.01	40.34	30.35	46.13	23.51	34.37			
	HEMBA1000024	16.68	21.57	43.98	24.42	20.7	22.54	19.03	14.92	22.2			
	HEMBA1000025	4.76	6.51	12.97	5.38	4.72	5.89	4.38	2.53	3.35			
10	HEMBA1000030	20.29	17.68	33.24	12.16	16.77	10.61	16.77	10.43	14.37			
	HEMBA1000036	14.36	14.21	18.16	2.88	10.75	5.3	7.22	6.69	3.23	*	**	-
	HEMBA1000037	11.11	16.21	12.93	4.27	8.42	4.3	6.83	2.67	4.79	*	*	-
	HEMBA1000039	8.22	15.6	26.39	9.92	10.66	6.88	12.98	7.81	13.65			
	HEMBA1000044	11.18	16.44	31.14	11.2	12.12	8.02	10.52	6.21	8.17			
	HEMBA1000048	8.23	14.79	19.17	7.61	10.2	7.2	15.14	3.23	4.66			
15	HEMBA1000050	12.03	11.84	30.06	14.4	13.92	13.02	11.35	6.38	12.59			
	HEMBA1000054	10.07	11.4	26.35	13.58	11.4	7.91	11.76	6.55	12.26			
	HEMBA1000055	458.88	310.46	850.13	127.99	485.16	208.47	362.27	384.53	141.27			
	HEMBA1000059	37.26	47.72	97.02	40.48	50.48	36.83	47.77	33.17	34.67			
	HEMBA1000072	84.93	93.08	177.56	168.05	149.4	148.43	84.23	55.56	75.03			
	HEMBA1000081	14.03	23.77	42.88	15.97	19.18	19.43	14.35	5.64	10.31			
20	HEMBA1000083	11.03	18.35	42.57	17.47	14.86	14.26	18.14	9.46	16.73			
	HEMBA1000089	12.54	18.95	30.4	15.64	14.81	12.6	22.7	8.13	18.39			
	HEMBA1000094	10.67	10.41	27.93	13.66	21.42	16.12	14.95	8.63	10.48			
	HEMBA1000097	6.26	10.91	14.76	6.71	9.5	4.54	15.21	6.91	3.77			
	HEMBA1000099	14.89	12.94	32.83	17.41	17.02	13.25	13.99	10.91	12			
	HEMBA1000103	79.17	84.17	154.39	129.48	133.3	142.91	117.04	80.44	93.95			
25	HEMBA1000106	13.4	15.97	19.47	7.19	4.59	5.52	5.12	3.42	4.98	**	**	-
	HEMBA1000113	4.8	10.67	20.57	6.45	7.88	9.01	9.02	3.11	6.76			
	HEMBA1000119	3.97	10.01	14.45	4.41	4.33	3.8	9.37	5.02	5.7			
	HEMBA1000133	88.66	47.28	101.21	23.18	118.13	70.04	134.7	41.76	81.57			
	HEMBA1000134	13.38	15.77	31.72	20.8	11.19	14.5	6.59	5.64	9.53			
	HEMBA1000136	34.17	26.36	68.3	67.42	34.73	49.05	25.95	21.33	26.27			
30	HEMBA1000141	7.91	11.34	20.35	10.92	8.05	11.46	11.73	5.94	9.7			
	HEMBA1000144	8.4	6.34	18.36	7.4	9.42	9.69	10.21	5.05	11.22			
	HEMBA1000147	3.5	3.8	22	5.61	6.52	8.28	7.06	1.83	3.83			
	HEMBA1000152	1.94	5.3	5.59	10.75	5.38	4.1	5.98	1.86	3.56			
	HEMBA1000154	7.08	8.64	11.01	9.8	10.82	8.45	9.52	5.05	11.09			
	HEMBA1000155	7.22	13.83	13.07	9.95	12.99	12.51	9.62	5.43	10.29			
35	HEMBA1000173	36.12	47.88	77.86	40.67	28.34	43.42	25.94	23.15	34.58			
	HEMBA1000175	4.3	6.41	7.7	6	3.69	3.11	3.98	4.45	4.04			
	HEMBA1000176	23.34	24.4	50.1	18.86	26.55	28.15	17.6	14.1	16.3			
	HEMBA1000196	1.73	5.17	9.68	5.19	3.63	4.66	4.31	1.03	3.4			
	HEMBA1000208	2.42	2.22	6.16	4.71	3	3.25	2.71	1.91	2.68			
	HEMBA1000209	4.99	7.73	19.16	5.65	5.94	7.5	3.72	1.31	4.4			
40	HEMBA1000212	4.66	5.69	7.88	3.69	6.97	3.08	4.71	1.75	3.95			
	HEMBA1000216	16.57	20.53	27.19	17.82	25.31	15.88	26.39	12.95	25.36			
	HEMBA1000217	25.38	37.33	52.96	16.88	15.01	12.09	9.3	23.54	15.2	*		-
	HEMBA1000218	31.37	54.57	73.09	37.79	38.52	34.06	37.08	24.06	36.49			
	HEMBA1000226	12.28	16.29	20.13	10.11	7.98	11.16	5.4	9.82	9.11	*		-
	HEMBA1000230	5.26	3.6	7.07	0.71	2.52	2.36	2.7	2.25	2.54	*	*	-
45	HEMBA1000240	5.06	6.06	9.84	4.39	4.5	5.46	2.75	2.24	3.3	*		-
	HEMBA1000244	7.48	12.78	15.39	11.19	10.98	11.37	6.63	4.57	10.35			
	HEMBA1000250	1.27	3.14	7.4	1.44	3.59	2.75	1.47	0.52	2.71			
	HEMBA1000258	10.17	16.3	24.98	15.18	17.62	13.97	14.16	8.49	16.99			
	HEMBA1000264	26.46	35.45	52.88	26.93	18.98	21.08	16.44	18.13	29.32			
	HEMBA1000266	12	14.67	15.28	5.33	5.8	4.87	9.31	5.79	7.42	**	*	-
50	HEMBA1000272	8.27	9.13	14.25	3.35	4.95	7.95	4.41	2.77	4.49	*		-
	HEMBA1000274	6.98	5.53	13.54	4.71	5.52	4.64	3.24	3.87	5.65			
	HEMBA1000276	5.09	3.14	11.53	3.19	3.43	4.43	2.41	0.94	4.11			
	HEMBA1000284	6.8	5.53	21.28	3.19	4.36	4.46	4.47	2.08	4.87			
	HEMBA1000307	8.46	14.9	24.21	13.5	12.6	8.38	8.3	4.98	7.88			
	HEMBA1000309	5.61	8.47	14.17	8.25	9.84	9.39	9.88	4.06	6.44			
55	HEMBA1000312	6.97	8.62	14.26	14.76	8.04	11.17	5.36	4.07	7.05			
	HEMBA1000317	7.52	4.64	7.99	3.4	3.92	3.39	4.7	5.98	4.31	*		-
	HEMBA1000318	6.93	8.7	14.09	8.17	6.23	6.97	7.59	3.11	9.29			

Table 380

	HEMBA1007051	8.21	9.64	21.16	10.38	7.94	9.6	8.59	4.82	5.46			
	HEMBA1007052	0.84	3.34	5.89	3.8	5.54	5	4.04	2.23	3.75			
5	HEMBA1007053	13.43	20.61	15.83	23.76	9.7	6.82	3.37	2.88	5.47	**	-	
	HEMBA1007057	6.84	7.67	12.49	5.42	9.68	14.27	10.77	4.02	8.83			
	HEMBA1007062	5.25	11.04	5.4	4.94	7.78	6.88	7.11	2.74	5.57			
	HEMBA1007063	19.82	23.76	40.56	22.16	18.37	20.33	18.69	15.25	20.25			
	HEMBA1007066	7.17	6.3	7.54	7.55	3.16	6.35	4.34	2.31	5.66	*	-	
10	HEMBA1007069	6.08	6.94	13.93	10.41	5.55	7.88	5.56	3.75	4.11			
	HEMBA1007073	5	7.14	13.19	9.06	6.96	8.85	10.68	5.02	4.92			
	HEMBA1007076	9.13	8.92	19.82	9.03	17.18	17.73	15.84	7.01	8.52			
	HEMBA1007078	47.57	62.26	92.7	36.29	60.73	51.72	32.07	18.35	34.97			
	HEMBA1007080	86	60.21	81.69	89.81	65.76	74.19	50.6	40.71	38.08	*	-	
	HEMBA1007084	10.6	15.68	20.31	13.17	16.74	11.58	13.37	8.47	13.17			
15	HEMBA1007085	27.29	31.32	26.53	13.81	11.35	9.85	7.06	8.85	9.17	**	**	-
	HEMBA1007087	17.6	23.13	29.36	17.3	13.84	15.36	6.49	22.47	10.72			
	HEMBA1007089	21.17	21.1	33.63	20.7	38.69	28.03	18.24	32.57	32.51			
	HEMBA1007095	201.88	116.22	315.3	326.58	303	223.56	75.5	112.08	127.01			
	HEMBA1007101	27.84	40.49	45.85	23.93	27.74	34.28	41.09	23.85	46.77			
20	HEMBA1007104	4.09	6	7.77	5.68	5.26	4.31	5.49	1.3	6.52			
	HEMBA1007106	21.73	33.28	23.78	16.4	24.34	20.37	10.72	5.09	4.25	**	-	
	HEMBA1007112	9.74	14.28	16.52	10.26	12.13	8.53	8.39	3.2	3.6	*	-	
	HEMBA1007113	21.85	25.07	43.45	24.93	19.55	20.96	26.57	21.39	27.54			
	HEMBA1007121	143.44	168.43	201.05	310.6	189.7	223.39	139.17	100.56	162.13			
	HEMBA1007129	6.41	8.28	18.59	3.42	2.84	6.23	5.66	3.45	4.93			
25	HEMBA1007147	12.43	15.07	27.38	13.28	12.69	11.69	10.77	9.49	8.81			
	HEMBA1007149	15.43	22.2	36.85	20.24	33.61	28.67	16.87	12.22	16.92			
	HEMBA1007151	6.66	4.91	10.19	4.89	4.76	3.64	3.37	0.61	3.29			
	HEMBA1007172	5.54	5.79	15.25	6.45	6.57	4.48	5.2	2.26	4.93			
	HEMBA1007174	5.55	6.74	11.4	7.96	7.31	3.31	8.36	1.4	3.1			
	HEMBA1007176	10.89	13.76	22.08	12.85	11.37	10.5	5.77	8.47	15.97			
30	HEMBA1007178	12.93	22.04	21.56	10.13	15.89	11.16	5.44	5.45	3.49	**	-	
	HEMBA1007185	12.08	13.89	28.24	5.12	12.57	13.87	7.15	7.65	10.67			
	HEMBA1007186	7.21	3.2	19.91	4.57	4.73	5.62	4.03	0.89	3.73			
	HEMBA1007194	17.48	18.75	30.8	17.47	18.26	18.1	18.81	10.34	19.89			
	HEMBA1007200	4.21	4.82	12.05	2.35	4.05	3.94	2.19	1.93	1.96			
	HEMBA1007203	17.41	23.17	30.14	16.76	21.31	8.75	11.68	7.36	6.25	*	-	
35	HEMBA1007206	25.5	42.75	37.89	30.71	35.28	24.97	38.96	21.71	36.83	*	**	-
	HEMBA1007224	24.39	40.17	34.49	15.91	19.51	9.53	6.52	8.53	10.32	*	**	-
	HEMBA1007226	89.72	73.23	106.85	134.34	88.67	116.09	67.3	41.82	79.65			
	HEMBA1007240	13.54	17.35	25.69	5.08	7.14	11.16	4.24	5.15	6.41	*	-	
	HEMBA1007241	5.35	7.7	9.62	3.24	3.12	3.91	4.59	2.56	3.21	*	*	-
	HEMBA1007242	4.41	5.32	8.31	2.05	4.24	3.88	2.68	0.75	5.03			
	HEMBA1007243	111.71	101.13	126.82	187.45	120.72	165.9	80.87	65.01	83.79	*	-	
40	HEMBA1007251	4.65	5.56	13.42	3.87	5.2	4.27	3.75	1.77	3.34			
	HEMBA1007256	7.71	17.43	18.8	11.41	15.37	11.45	13.72	5.68	6.36			
	HEMBA1007267	27.14	33.68	57.79	28.23	30.36	27.93	22.24	21.62	27.1			
	HEMBA1007273	5.82	10.89	12.53	5.12	4.08	4.32	3.9	1.99	5.52			
	HEMBA1007279	8.03	6.83	14.11	6.96	6.78	5.74	5.78	2.58	3.49			
	HEMBA1007281	4.41	6.17	11.59	1.95	1.64	1.99	3.42	2.15	2.75			
45	HEMBA1007283	10.13	14.5	18.59	11.27	10.46	9.5	8.86	8.35	11.65			
	HEMBA1007288	10	10.12	23.07	7.33	8.9	6.31	9.07	7.22	9.19			
	HEMBA1007291	3.94	6.53	14.21	2.35	5.74	4.15	6.79	3.29	5.42			
	HEMBA1007299	60.36	57.45	88.44	47.01	83.94	54.92	41.98	17.99	24.76	*	-	
	HEMBA1007300	10.43	10.11	14.15	17.47	5.8	7.51	10.92	2.6	3.93			
	HEMBA1007301	10.28	11.15	10.5	7.08	5.07	5.8	3.91	4.02	3.67	**	**	-
50	HEMBA1007319	7.15	12.6	10.93	3.57	6.52	4.89	3.16	3.33	2.24	*	*	-
	HEMBA1007320	6.22	6.62	11.18	5.17	5.74	4.76	5.15	2.1	2.94			
	HEMBA1007322	281.14	504.93	265.82	479.23	433.13	233.63	384.76	246.29	107.4			
	HEMBA1007323	3.87	5.98	13.69	3.16	3.64	1.67	4.69	0.43	1.85			
	HEMBA1007326	33.32	37.02	90.88	53.8	48.19	37.22	52.37	23.73	38.43			
	HEMBA1007327	14.51	15.7	50.08	17.82	23.55	19.83	20.48	9.78	17.51			
55	HEMBA1007332	15.93	13.1	23.53	14.44	19.66	12.93	14.13	7.11	7.2			
	HEMBA1007341	8.96	9.53	11.63	7.92	6.35	6.27	7.48	4.06	5.93	*	*	-
	HEMBA1007342	5.2	5.9	6.99	1.75	3.52	2.76	4.48	1.49	2.47	*	*	-

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	HENBA1006754	10.3	5.77	11.65	6.17	5.72	5.75	6.13	5.01	8.39			
	HENBA1006758	7.27	4.9	12.62	10.94	5.09	5.73	5.5	2.15	4.84			
5	HENBA1006767	5.55	6.53	9.74	2.73	5.09	3.33	3.97	3.29	3.02	*		-
	HENBA1006770	10.91	18.24	23.01	9.34	17.17	9.49	6.46	5.37	4.52	*		-
	HENBA1006779	21.14	29.08	46.61	19.18	32.07	21.74	27.02	14.11	23.35			
	HENBA1006780	31.66	41.91	71.99	39.02	25.6	33.44	20.75	21.87	35.28			
	HENBA1006789	11.15	8.92	10.14	6.24	5.48	5.24	3.29	3.78	6.02	**	**	-
	HENBA1006795	13.29	18.12	23.2	10.87	8.5	11.23	14.12	7.4	15.69			
10	HENBA1006796	8.15	7.65	12.79	2.99	3.39	3.14	3.97	0.85	2.9	*	*	-
	HENBA1006805	10.17	13.14	20.22	7.25	5.84	4.86	8.13	5.47	7.83	*		-
	HENBA1006807	358.41	425.1	450.14	425.56	524.37	388.31	479.65	356.88	266.31			
	HENBA1006813	3.32	4.75	8.38	0.01	2.55	1.47	2.36	0.6	1.87			
	HENBA1006819	11.44	14.72	25.12	8.56	18.43	11.48	10.78	2.32	9.36			
15	HENBA1006821	15.76	14.89	42.37	23.12	22.95	19.27	10.56	11	21.11			
	HENBA1006824	31.97	30.86	49.16	24.26	23.62	19.71	31.05	26.34	34.02			
	HENBA1006832	259.36	308.26	291.59	339.91	237.21	321.61	326.22	280.59	176.89			
	HENBA1006834	46.64	65.27	67.81	62.09	43.11	55.61	73	47.8	42.25			
	HENBA1006835	8.46	14.27	21.58	9.55	10.78	10.45	9.6	6.1	8.18			
	HENBA1006843	18.15	18.91	28.2	12.23	21.78	20.94	21.53	7.17	5.58			
20	HENBA1006849	26.18	26.08	44.03	23.34	31.15	21.57	33.36	14.9	24.69			
	HENBA1006850	16.77	33.56	37.75	10.59	18.63	8	9.03	3.5	6.56	*		-
	HENBA1006861	14.67	13.53	34.61	20.34	16.15	19.04	12.38	6.63	17.89			
	HENBA1006865	39.2	38.77	61.65	71.58	60.52	57.06	33.27	28.86	36.65			
	HENBA1006867	18.5	15.53	22.34	9.22	8.88	9.34	9.42	6.3	3	**	**	-
	HENBA1006873	17.71	23.93	26.16	14.12	11.71	14.57	15.58	7.18	17.11	*	*	-
25	HENBA1006877	7.4	13.49	15.12	6.73	6.6	3.67	5.7	2.95	4.47	*		-
	HENBA1006878	10.48	14.11	25.64	11.23	13.75	11.88	9.01	2.84	7.35			
	HENBA1006879	12.94	16.27	24.73	12.26	12.67	8.96	12.1	3.4	9.75			
	HENBA1006884	16.51	27.4	33.59	6.17	20.26	51.39	18.62	5.59	4.55			
	HENBA1006885	29.88	29.68	41.54	20.77	25.78	20.37	13.71	10.53	21.42	*		-
	HENBA1006886	94.54	114.96	86.8	8.15	45.42	40.45	16.06	32.92	14.53	**	**	-
30	HENBA1006889	11.07	12.36	15.21	4.3	6.85	5.29	4.87	2.48	1.64	**	**	-
	HENBA1006896	72.84	75.71	95.1	25.42	32.71	26.4	32.9	16.88	9.13	**	**	-
	HENBA1006900	25.47	32.8	47.12	42.47	30.76	32.5	22.96	11.97	13.31			
	HENBA1006902	4.95	12.4	20.5	3.65	6.21	3.85	6.17	0.96	2.8			
	HENBA1006912	17.87	28.61	49.99	20.94	22.1	17.02	26.3	12.33	21.55			
	HENBA1006914	76.35	138.05	152.36	81.86	82.01	60.45	85.41	43.28	74.2			
35	HENBA1006916	15.23	15.18	19.35	6.9	10.02	9.36	4.27	4.91	5.74	**	**	-
	HENBA1006921	5.69	9.45	11.27	4.84	7.39	7.2	5.96	4.4	4.47			
	HENBA1006926	14.39	14.83	25.68	4.9	10.25	8.6	7.17	9.16	6.2	*		-
	HENBA1006927	8.1	6.16	12.12	3.55	6.9	3.55	4.27	4.44	3.56			
	HENBA1006929	20.15	20.6	39.98	22.07	23.94	22.4	25.87	18.49	21.97			
	HENBA1006936	9.54	10.99	23.15	1.82	9.22	9.34	13.47	4.19	5.58			
40	HENBA1006938	3.12	8.25	12.19	4.54	5.27	8.96	3.85	0.26	1.77			
	HENBA1006941	38.07	33.9	81.07	14.16	37.03	19.11	37.54	10.57	9.16			
	HENBA1006942	17.04	18.44	31.05	13.54	11.67	15.34	9.95	7.49	11.08			
	HENBA1006945	67.94	90.41	117.96	45.68	44.82	62.84	58.07	38.55	55.06			
	HENBA1006949	5.04	6.37	14.93	11.38	7.05	7.71	7.34	3.55	6.74			
	HENBA1006952	4.19	5.26	9.54	3.87	5.23	3.34	4.22	1.4	1.92			
45	HENBA1006950	34.21	40.14	82.69	40.95	44.04	37.38	52.24	36.27	39.73			
	HENBA1006973	11.14	12.47	24.84	14.07	12.49	8.94	10.47	5.46	6.01			
	HENBA1006974	9.69	21.72	26.59	5.58	10.54	8	10.23	6.86	3.65			
	HENBA1006976	5.49	4.82	13.28	5.67	2.91	2.68	2.79	1.22	3.42			
	HENBA1006989	5.64	5.97	15.87	14.35	9.67	15.25	5.09	3.23	6.48			
	HENBA1006993	22.88	33.04	62.59	43.82	36.96	35.47	29.4	20.95	25.28			
50	HENBA1006996	6.77	6.15	10.45	9.29	8.92	7.56	4.85	2.62	3.17	*		-
	HENBA1007001	13.6	18.68	46.24	26.01	27.9	24.22	17.93	11.6	13.83			
	HENBA1007002	38.2	51.66	86.51	157.46	90.61	115.16	37.83	43.86	26.69			
	HENBA1007013	4.23	5.6	9.85	5.49	4.83	5.04	4.32	1.34	5.39			
	HENBA1007016	13.73	18.8	46.37	32.25	24.75	25.55	26.93	13.32	18.08			
	HENBA1007017	1.76	2.36	7.34	6.45	3.5	5.8	4.75	0.45	2.16			
55	HENBA1007018	12.78	19.44	26.15	6.88	7.17	10.67	5.95	5.18	4.75	*		-
	HENBA1007044	4.27	3.89	8.13	7.26	4.08	4.08	4.49	2.95	4.25			
	HENBA1007045	3.76	6.62	12.6	6.66	6.82	7.16	7.91	2.23	5.58			

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	HENBA1006483	10.38	11.79	17.97	10.54	8.06	5.41	9.39	6.26	10.8				
	HENBA1006485	14.11	21.34	22.05	9.66	9.42	6.66	9.41	4.35	6.34	*	*	-	-
5	HENBA1006486	25.6	38.65	46.49	15.25	21.42	18.99	28.29	12.46	17.42	*		-	-
	HENBA1006489	4.69	5.66	11.99	2.16	3.64	2.6	4.74	2.19	2.49		*	-	-
	HENBA1006492	21.26	21.32	31.79	8.11	21.1	11.99	16.25	6.33	8.98		*	-	-
	HENBA1006494	3.73	5.82	13.17	4.37	5.86	3.22	4.57	1.78	4.99		*	-	-
	HENBA1006497	9.48	5.39	9.72	6.41	4.7	6.22	4.84	2.41	5.11		*	-	-
	HENBA1006501	70.71	74.3	109	137.89	121.18	132.43	56.47	47.55	54.22	*	*	+	-
10	HENBA1006502	26	37.84	37.28	17.65	19.66	21.79	14.53	13.47	14.09	*	**	-	-
	HENBA1006507	74.35	71.35	93.56	76.72	76.19	89.01	54.44	45.62	81.01			-	-
	HENBA1006517	10.15	13.34	25.2	8.05	9.91	8.2	9.98	8.78	9.53			-	-
	HENBA1006521	4.71	7.89	15.55	2.17	4.99	3.48	4.73	1.47	4.08		*	-	-
	HENBA1006529	20.44	40.14	54.85	11.1	18.97	9.04	8.44	3.17	5.16		*	-	-
	HENBA1006530	2.11	3.93	16.06	3.3	3.94	2.97	2.36	0.8	2.85		*	-	-
15	HENBA1006535	4.76	6.35	8.43	4.93	4.07	4.99	1.52	1.3	3.36		*	-	-
	HENBA1006536	7.73	9.35	15.74	6.97	6.8	8.19	6.06	4.78	7.71		*	-	-
	HENBA1006540	4.89	5.07	10.1	3.8	4.1	3.21	4.5	0.57	2.57		*	-	-
	HENBA1006544	9.55	9.44	12.65	5.51	6.6	3.64	6.81	6.28	5.66	*	*	-	-
	HENBA1006546	13.83	18.34	35.37	18.49	13.07	8.02	20.42	13.65	12.24			-	-
	HENBA1006549	3.69	4.64	10.47	3.5	3.22	3.38	4.38	1.91	2.6			-	-
20	HENBA1006559	5.63	7.78	15.93	4.1	5.24	3.33	4.33	1.27	1.6			-	-
	HENBA1006562	4.46	7.73	13.41	4.81	4.36	7.66	6.04	1.8	5.11			-	-
	HENBA1006566	1.97	2.79	6.04	4.85	3.75	3.09	2.63	0.98	3.24			-	-
	HENBA1006569	9.03	13.38	17.77	8.08	5.02	11.82	6.18	3.56	5.28	*	*	-	-
	HENBA1006572	4.01	5.33	6.28	3.85	4.61	4.82	3.06	1.11	1.97	*	*	-	-
	HENBA1006579	48.81	48.67	106.01	77.85	58.85	72	32.03	38.94	29.18			-	-
25	HENBA1006583	21.83	28.38	52.62	37.84	32.54	33.86	32.6	18.06	30.33			-	-
	HENBA1006595	9.39	11.2	24.82	12.48	12.13	12.39	17.97	4.53	9.41			-	-
	HENBA1006597	10.74	17.9	34.13	14.71	18.52	13.99	17.64	9.01	20.29			-	-
	HENBA1006606	9.14	17.8	25.4	12.84	13.75	11.34	17.49	5.73	12.78			-	-
	HENBA1006612	16.16	16.45	27.36	16.16	10.37	13.84	7.68	7.75	9.58	*	*	-	-
	HENBA1006617	8.08	9.55	19.24	13.65	9.22	10.71	8.25	4.71	9.91			-	-
30	HENBA1006624	62.58	66.3	115.43	55.71	47.06	62.71	45.87	24.82	44.31			-	-
	HENBA1006631	23.12	25.87	43.15	23.48	22.27	20.06	24.55	15.53	27.45			-	-
	HENBA1006635	7.91	10.88	14.85	9.02	10.78	9.39	11.43	5.66	11.05			-	-
	HENBA1006639	10.04	12.39	17.07	18.66	19.19	12.58	8.27	4.47	14.67			-	-
	HENBA1006643	1.84	4.34	3.2	4.98	3.53	3.26	2.93	0.82	1.75			-	-
	HENBA1006648	15.53	17.03	27.49	19.15	15.29	16.67	5.29	8.82	12.26			-	-
35	HENBA1006652	26.71	34.75	50.3	32.97	21.76	18.68	16.24	13.51	23.67			-	-
	HENBA1006653	7.56	6.61	17.15	7.18	4.77	7.22	2.33	2.95	3.74			-	-
	HENBA1006658	19.91	14.88	49.21	18.71	18.4	21.38	11.54	8.35	7.4			-	-
	HENBA1006659	58.61	60.76	125.49	159.5	103.52	108.54	57.82	49.19	67.43			-	-
	HENBA1006665	2.46	4.7	11.42	4.57	4.2	3.45	4.37	1.96	4.61			-	-
	HENBA1006666	3.78	4.89	6.54	3.83	3.74	5.58	5.07	2.85	4.33			-	-
40	HENBA1006671	29.88	46.04	31.81	11.03	10.78	7.48	20.4	8.57	17.2	**	*	-	-
	HENBA1006674	17.95	21.64	48.41	21.77	17.05	22.65	20.34	20.43	18.22		*	-	-
	HENBA1006676	8.33	14.71	17.2	11.8	6.88	5.97	6.74	4.48	6.11		*	-	-
	HENBA1006682	5.28	7.81	6.48	5.22	3.34	3.44	5.44	2.35	3.91		*	-	-
	HENBA1006688	8.06	7.99	15.76	5.58	7.11	7.53	6.82	4.86	5.53		*	-	-
	HENBA1006695	11.68	16.81	23.72	8.98	10.42	13.13	10.55	9	11.41		*	-	-
45	HENBA1006696	12.25	15.75	12.83	8.51	9.6	6.76	4.49	2.19	4.51	*	**	-	-
	HENBA1006702	6.88	9.14	9.5	5.54	6.34	5.86	4.33	1.73	4.63	*	*	-	-
	HENBA1006707	12.13	15.02	19.11	19.48	16.83	14.64	14.25	5.63	13.01			-	-
	HENBA1006708	7.17	12.19	21.37	12.13	8.06	8.49	3.77	4.26	7.59			-	-
	HENBA1006709	13.09	19.04	28.5	14.85	14.05	15.86	11.87	11.01	16.42			-	-
	HENBA1006717	5.25	5.26	12.53	4.74	4.01	3.68	7.38	2.38	6.67			-	-
50	HENBA1006724	12.11	8.3	19.78	5.53	10.32	5.8	9.44	6.24	6.16			-	-
	HENBA1006731	7.48	7.6	18.92	6.67	8.76	6.75	5.19	3.77	3.96			-	-
	HENBA1006737	6.06	6.19	10.33	4.37	4.14	4.69	4.38	0.91	4.4			-	-
	HENBA1006742	8.98	8.7	24.13	7.5	9.95	8.07	10.84	6.04	17.08		*	-	-
	HENBA1006743	14.91	22.66	22.54	17.15	17.28	14.05	13.93	8.87	7.89		*	-	-
	HENBA1006744	17.72	19.61	39.58	28.17	20.27	23.33	18.16	15.95	23.07		*	-	-
55	HENBA1006749	4.84	4.51	4.65	3.84	2.79	5.06	3.99	2.53	5.17		*	-	-
	HENBA1006752	74.49	114.01	106.5	80.8	62.77	63.46	56.97	93.88	63.11		*	-	-

Table 377

	HENBA1006226	60.66	89.4	128.3	98.59	21.45	56.73	34.83	42.34	47.3		
	HENBA1006235	7.29	6.42	9.51	4.4	5.83	6.34	4	1.9	2.78	*	-
5	HENBA1006248	8.59	14.3	12.76	17.84	10.1	10.62	14.29	11.1	13.06		
	HENBA1006251	15.98	24.11	24.67	22.82	19.08	21.09	16.03	3.32	8.44		
	HENBA1006252	6.29	11.35	24.58	17.83	5.75	7.22	9.32	6.16	9.75		
	HENBA1006253	6.65	9.14	20.63	14.78	17.19	22.49	8.79	1.77	6.4		
	HENBA1006259	11.81	21.15	28.91	16.9	10.94	16.19	15.68	14.83	17.05		
10	HENBA1006261	25.45	37.82	48.94	33.85	29.71	14.07	18.76	15.97	24.68		
	HENBA1006268	4.53	5.62	11.14	5.03	3.83	5.91	4.8	2.63	2.52		
	HENBA1006271	27.87	48.98	84.46	39.21	39.6	40.82	30.74	14.23	28.33		
	HENBA1006272	3.48	5.09	12.53	3.82	7.27	10.33	8.77	2.68	5.16		
	HENBA1006273	4.26	5.62	9.16	7.22	6.74	6.25	4.33	1.91	3.78		
	HENBA1006276	8	7.56	13.01	10.92	8.88	11.33	8.59	5.55	8.16		
	HENBA1006278	3.56	6.86	4.87	4.56	7.03	4.18	3.68	1.91	2.81		
15	HENBA1006283	21.61	22.15	38.14	17.11	13.97	13.64	11.04	8.54	8.53	*	-
	HENBA1006284	9.77	12.43	9.21	7.3	6.48	8.66	4.11	18.95	4.13		
	HENBA1006291	7.57	5.98	8.47	4.12	5.97	5.86	4.51	2.35	4.32	*	-
	HENBA1006292	28.37	23.25	41.05	49.83	29.24	37.1	21.69	20.86	21.95	*	-
	HENBA1006293	3.75	5.49	8.43	3.08	2.38	10.01	0.91	2.39	2.11	*	-
20	HENBA1006299	20.83	28.58	31.92	37.12	28.94	29.71	18.96	13.53	20.58		
	HENBA1006309	12.21	13.85	16.63	10.12	12.48	15.92	14.59	3.8	13.91		
	HENBA1006310	10.98	14.98	14.35	11.71	17.38	10.36	10.75	8.47	4.15		
	HENBA1006311	7.24	7.64	16.35	13.11	5.74	8	7.15	4.03	4.85		
	HENBA1006313	3.22	8.11	10.2	4.45	3.61	3.07	2.82	2.44	2.58		
	HENBA1006316	7.65	12.92	15.52	10.38	5.11	11.89	7.9	6.94	9.74		
25	HENBA1006328	21.63	21.11	38.33	32.37	21.64	26.1	17.68	13.27	19.36		
	HENBA1006334	3.37	3.68	14.62	4.58	3.28	2.72	2.89	4.86	1.73		
	HENBA1006335	17.68	32.85	46.47	23.14	32.25	24.21	26.2	15.13	17.28		
	HENBA1006344	55.44	64.87	82.83	51.59	45.13	40.79	32.94	23.43	27.33	**	-
	HENBA1006347	7.7	7.61	11.32	7.96	6.93	5.21	8.91	1.81	2.49		
	HENBA1006349	5.12	7.39	13.08	11.96	7.92	10.21	7.73	3.41	8.88		
30	HENBA1006352	4.71	5.44	5.8	3.17	4.05	6.34	3.53	2.18	6.11		
	HENBA1006357	21.84	30.15	32.51	16.24	19.4	20.03	17.68	17.67	25.75	*	-
	HENBA1006358	9.29	7.19	14.9	8.1	8.4	7.37	7.12	3.93	6.97		
	HENBA1006359	5.5	5.93	13.36	4.55	6.66	5.77	4.91	2.59	3.15		
	HENBA1006360	4.46	4.17	9.74	2.49	4.18	4.08	2.89	1.16	2.66		
	HENBA1006364	4.83	5.25	14.76	2.78	4.9	3.21	2.71	2.21	3.79		
35	HENBA1006377	80.85	114.32	105.84	75.76	102.36	58.33	93.99	60.04	78.87		
	HENBA1006380	20.03	20.23	49.46	23.06	19.66	23.19	14.68	15.28	25.75		
	HENBA1006381	21.8	26.78	34.04	28.98	18.12	16.56	28.76	15.54	19.65		
	HENBA1006385	26.15	27.47	48.23	21.69	21.93	26.5	17.06	11.15	26.66		
	HENBA1006390	66.76	79.54	86.03	47.23	63.78	61.2	79.04	51.69	65.97		
	HENBA1006391	9.83	7.54	21.2	5.37	10.85	8.66	7.1	4.25	6.66		
40	HENBA1006398	7.74	10.16	17.94	6.15	8.15	10.18	5.31	3.87	6.36		
	HENBA1006405	17.21	30.3	36.97	11.67	17.28	9.3	9.52	5.99	10.52	*	-
	HENBA1006410	12.29	24.19	20.82	11.01	22.54	19.08	10.32	24.08	7.5		
	HENBA1006416	13.77	16.34	35.1	20.2	19.08	14.96	13.83	12.25	17.9		
	HENBA1006418	14.88	14.23	12.55	8.74	10.37	7.82	5.57	12.05	3.94	**	-
	HENBA1006419	30.32	26.38	45.23	21.57	20.76	20.5	35.29	23.16	35.95		
45	HENBA1006421	6.94	7.3	14.62	4.48	5.03	4.44	5.76	2.74	5.79		
	HENBA1006424	6.59	5.16	10.86	2.39	3.13	4.03	3.8	3.29	5.21		
	HENBA1006426	16.94	21.77	35.51	13.82	15.6	11.92	15.06	10.89	16.55		
	HENBA1006430	21.51	23.1	52.14	21.8	26.01	14.55	25.55	15.62	24.71		
	HENBA1006438	7.39	9.88	27.72	9.7	12.86	10.02	7.77	3.89	8.06		
	HENBA1006445	5.97	4.62	11.24	6.9	4.18	5.84	3.07	2.65	7.57		
50	HENBA1006446	10.44	29.9	25.23	12.03	4.68	7.84	2.86	8.87	4.65		
	HENBA1006456	20.27	20.9	27.71	19.85	19.2	23.69	16.72	10.9	15.13	*	-
	HENBA1006461	10.85	13.87	19.86	8.91	8.16	8.85	10.75	7.35	8.75		
	HENBA1006467	2.88	4.33	9.77	4.94	3.17	2.66	4.06	1.76	2.14		
	HENBA1006470	34.3	38.7	56.67	41.16	36.04	29.19	28.2	24.42	29.45		
	HENBA1006471	8.92	14.85	23.43	12.81	12.27	40.2	16.21	5.14	6.51		
55	HENBA1006474	80.88	55.42	113.87	93.82	77.51	77.92	59.58	47.32	61.33		
	HENBA1006476	124.18	125.67	311.61	282.54	252.79	111.47	88.12	112.31	128.83		
	HENBA1006482	750.18	800.51	717.99	555.6	740.67	596.85	785.76	753.48	372.51		

Table 376

	HEMBA1005907	11.51	24.95	21.68	5.18	12.96	7.51	14.52	20.83	32.4		
	HEMBA1005909	12.54	10.65	15.16	9.37	9.7	8.92	5.33	5.2	5.61	**	-
5	HEMBA1005911	4.25	2.94	8.69	4.53	4.07	3.47	3.19	0.81	2.14		
	HEMBA1005912	35.05	30.41	48.44	55.56	35.89	54.16	35.55	16.99	36.27		
	HEMBA1005913	8.07	12.66	17.16	3.35	11.55	8.12	4.56	4.08	5.57	*	-
	HEMBA1005921	8.39	17.95	29.27	13.5	18.47	18.42	20.02	11.79	13.13		
	HEMBA1005922	10.23	12.69	22.71	8.62	19.47	16.43	10.1	8.02	9.98		
10	HEMBA1005929	14.03	14.99	35	19.43	15.07	16.06	12.42	10.95	15.92		
	HEMBA1005931	6.92	7.45	8.96	7.96	7.11	4.38	5.8	5.53	5.56	*	-
	HEMBA1005934	9.15	11.61	14.52	6.47	9.42	7.17	6.06	6.07	3.78	*	-
	HEMBA1005945	13.18	17.17	24.76	14.61	10.38	10.87	14.78	9.7	14.04		
	HEMBA1005962	8.03	8.91	21.61	8.95	9.17	9.16	8.64	3.56	9.28		
	HEMBA1005963	7.03	8.04	15.69	2.68	5.58	6.69	5.62	2.88	7		
15	HEMBA1005990	1.77	6.11	10.33	2.7	4.79	2.9	2.4	1.78	3.08		
	HEMBA1005991	64.98	101.38	91.59	21.08	89.91	49.01	45.05	21.27	28.13	*	-
	HEMBA1005999	10.77	8.18	14.85	9.74	9.96	11.62	9.75	7.35	11.3		
	HEMBA1006002	4.73	10.03	9.92	4.44	5.47	3.19	2.29	3.3	1.96	*	-
	HEMBA1006005	11.33	12.26	21.89	15.72	14.29	10.03	11.77	7.68	11.25		
	HEMBA1006011	7.74	6.02	9.99	4.17	3.85	3.65	4.89	1.78	2.14	*	*
20	HEMBA1006013	15.75	24.43	27.02	14.33	11.65	10.41	12.44	6.76	8.78	*	*
	HEMBA1006016	12.67	16.72	32.68	4.67	8.33	5.18	11.34	2.62	5.66		
	HEMBA1006019	39.35	43.51	94.43	41.01	47.95	33.8	48.65	20.74	35.48		
	HEMBA1006021	100.18	64.95	186.84	161.47	154.88	173.68	76.07	51.55	67.34		
	HEMBA1006022	4.41	4.65	8.97	5.98	4.93	5.29	5.93	2.38	6.8		
	HEMBA1006031	54.5	55.09	63.41	23.75	30.34	33.23	14.55	24.74	14.18	**	**
25	HEMBA1006035	26.81	26.57	28.41	13.64	17.75	10.2	23.49	16.85	24.27	**	-
	HEMBA1006036	30.41	35.58	51.17	28.85	21.84	21.47	22.61	12.69	16.92	*	-
	HEMBA1006042	14.78	15.89	30.2	21.62	13.62	17.35	12.05	7.06	10.83		
	HEMBA1006044	6.37	6.38	15.67	2.89	5.13	3.3	5.6	2.77	4.16		
	HEMBA1006045	5.05	5.98	16.26	5.37	6.51	3.98	4.26	3.34	4.33		
	HEMBA1006048	5.66	8.21	17.12	5.57	7.7	7.23	7.81	2.07	6.93		
30	HEMBA1006053	9.91	12.14	22.17	11.19	10.11	10.25	6.6	4.29	8.53		
	HEMBA1006055	18.11	16.89	29.58	15.19	13.98	12.78	12.35	11.58	12.66		
	HEMBA1006058	12.02	10.46	14.6	9.27	7.95	6.43	9.22	7.8	6.16	*	*
	HEMBA1006063	115.82	107.87	141.14	161.05	129.88	130.01	94.43	64.84	83.98	*	-
	HEMBA1006067	14.46	16.19	18.97	8.28	13.91	5.01	10.46	4.28	4.98	*	-
	HEMBA1006081	3.95	8.3	15.7	2.09	6.94	3.38	3.92	1.35	3.37		
35	HEMBA1006089	4.09	6.69	17.6	2.93	4.85	1.66	3.54	1.57	3.85		
	HEMBA1006090	16.61	33.98	39.21	14.19	23.81	18.25	18.94	5.99	5.67		
	HEMBA1006091	6.4	6.94	10.5	5.04	5.34	5.02	4.02	3.3	3.85	*	-
	HEMBA1006093	24.23	35.95	33.59	5.11	7.76	13.97	5.79	10.1	7.62	**	**
	HEMBA1006099	23.7	22.12	46.77	18.77	21.87	13.39	21.06	17.59	21.54		
	HEMBA1006100	15.19	17.46	28.4	13.14	14.55	10.6	18.9	11.15	17.08		
40	HEMBA1006108	9.23	5.57	12.26	4.66	7.02	4.99	4.59	2.55	4.37		
	HEMBA1006114	6.24	4.2	16.8	2.53	7.16	4.48	5.37	0.85	3.06		
	HEMBA1006121	4.96	7.47	17.94	3.37	6.22	2.88	5.11	1.02	2.9		
	HEMBA1006124	23.37	10.77	43.63	10.18	13.13	12.16	9.91	5.78	5.28		
	HEMBA1006125	5.3	5.09	9.35	4.55	4.9	6.58	2.71	1.37	4.21		
45	HEMBA1006130	81.41	100.18	218.91	239.88	178.96	190.17	100.6	68.45	89.08		
	HEMBA1006138	16.78	21.47	42.54	20.91	18.85	14.71	18.67	11.93	13.92		
	HEMBA1006142	6.25	5.14	9.74	3.49	5.53	4.08	2.75	2.92	2.44	*	-
	HEMBA1006150	20.94	16.77	47.47	16.05	12.94	20.13	22.73	16.22	18.83		
	HEMBA1006151	3.81	3.62	12.07	4.06	2.71	3.79	4.85	1.6	3.25		
	HEMBA1006155	51.87	36.97	81.85	14.8	33.19	24.55	29.86	21.38	8.77		
	HEMBA1006158	7.8	6.18	16.28	4	7.84	5.44	10.27	1.55	3.09		
50	HEMBA1006164	7.41	4.45	10.51	11.19	6.62	6.2	5.41	2.39	2.71		
	HEMBA1006171	11.38	13.76	11.95	10	9.74	12.84	7.5	4.2	3.45	**	-
	HEMBA1006173	5.77	5.17	5.55	5.75	11.01	7.35	5.28	2.14	3.83		
	HEMBA1006176	10.21	6.39	20.68	12.36	13.62	14.28	6.26	8.81	6.76		
	HEMBA1006182	6.52	8.53	12.82	8.56	9.55	11.37	5.71	3.32	5.23		
	HEMBA1006197	4.36	5.59	8.45	4.07	4.78	4.63	3.18	1.47	2.61	*	-
55	HEMBA1006198	6.83	8.29	18.43	9.27	9.93	14.63	6.95	4.93	6.04		
	HEMBA1006213	46.5	59.16	87.76	45.37	52.9	41.61	64.69	38.82	49.81		
	HEMBA1006217	77.53	104.19	150.86	44.45	56.82	81.63	56.2	53.63	65.92		

Table 375

	HENBA1005576	3.76	5.34	5.63	3.54	6.15	2.3	3.82	3.83	3.35			
	HENBA1005577	3.45	4.03	5.12	2.3	2.95	1.9	2.63	1.06	1.56	*	*	-
5	HENBA1005581	17.34	15.11	23.24	3.89	10.6	7.1	4.64	4.62	3.74	*	**	-
	HENBA1005582	240.9	356.29	365.46	317.4	608.67	346.15	243.13	155.96	316.09			
	HENBA1005583	56.28	75.41	54.74	36.66	53.45	30.25	44.93	12.37	20.67	*	*	-
	HENBA1005588	48.19	61.31	71.13	27.32	36.77	18.93	99.81	56.7	45.14	*	*	-
	HENBA1005593	8.46	13.58	28.24	14.33	10.97	7.86	11.3	3.59	6.67			
	HENBA1005595	5.4	4.89	14.77	3.77	3.53	5.64	3.73	2.81	4.24			
10	HENBA1005597	4.61	6.07	5.65	4.04	20.76	7.93	6.41	5.62	16.69			
	HENBA1005606	10.57	19.2	23.77	3.86	10.94	4.17	5.54	1.39	3.32	*	*	-
	HENBA1005609	15.17	18.03	17.77	3.19	7.05	3.09	3.33	3.84	3.16	**	**	-
	HENBA1005616	3.29	3.65	8.89	1.88	4.28	2.06	3.75	2.04	3.26			
	HENBA1005621	2.8	5.27	13.13	1.63	2.71	2.81	4.45	1.3	2.19			
	HENBA1005627	26.17	45.68	60.66	11.35	32.13	15.73	25.61	10.31	7.56			
15	HENBA1005628	4.16	6.62	18.11	5.71	6.97	5.93	6.45	1.4	3.09			
	HENBA1005631	2.97	2.99	6.58	2.7	2.52	4.68	1.93	1.12	1.83			
	HENBA1005632	8.75	7.76	13.3	5.6	4.94	4.75	6.21	2.5	5.1	*	*	-
	HENBA1005634	13.26	27.71	36.18	8.65	7.52	11.69	10.2	6.61	5.28			
	HENBA1005662	5.49	5.63	8.59	1.07	4.9	3.69	4.47	3.62	2.83			
	HENBA1005666	4.1	8.92	17.42	4.51	5.05	7.36	8.37	3.56	4.18			
20	HENBA1005670	19.65	25.57	58.84	10.77	25.74	28.55	25.33	19.21	13.78			
	HENBA1005671	3.25	3.78	12.05	4.13	4.03	2.36	2.63	0.58	1.59			
	HENBA1005679	30.48	23.92	60.53	11.99	24.03	21.81	34.62	12.99	24.28			
	HENBA1005680	5.31	5.25	11.55	7.98	6.67	8.25	4.31	2.95	4.88			
	HENBA1005685	10.54	10.31	19.75	6.25	14.08	11.58	12.25	7.49	2.77			
	HENBA1005698	3.05	4.09	5.98	5.75	3.35	4.26	2.89	1.71	1.85			
25	HENBA1005699	81.55	73.56	131.24	173.55	121.01	131.74	58.39	29.63	52.24			
	HENBA1005703	8.7	6.7	19.33	15.32	12.15	10.29	11.29	5.53	9.29			
	HENBA1005705	10.68	16.62	28.52	12.65	15.48	9.89	13.24	7.86	8.73			
	HENBA1005712	14.41	9.07	21.89	4.52	12.06	8.24	12.54	7.16	9.37			
	HENBA1005717	9.58	15.11	28.28	15.34	17.37	16.39	10.48	4.92	9.53			
	HENBA1005718	14.72	17.46	32.28	23.31	13	19.82	12.33	8.89	13.45			
30	HENBA1005721	7.61	8.19	16.43	10.15	9	9.16	5.24	3.31	7.52			
	HENBA1005722	8.96	8.38	15.56	10.88	5.15	8.22	6.16	2.61	4.2			
	HENBA1005724	3.21	3.63	9.19	3.08	3.41	4.96	4.32	2.79	3.41			
	HENBA1005732	26.35	14.39	37.65	46.13	33.11	35.41	27.65	13.58	24.67			
	HENBA1005737	10.7	10.58	19.74	12.96	11.14	14.53	14.77	5.18	11.46			
	HENBA1005742	24.19	21.22	31.29	17.65	22.38	19.25	19.89	11.57	20.16			
35	HENBA1005746	16.18	18.76	28.18	19.8	21.24	20.22	17.64	16.51	21.84			
	HENBA1005747	4.59	6.97	9.33	5.84	4.66	4.23	3.4	2.42	2.9	*	*	-
	HENBA1005749	4.06	4.65	5.89	4.59	4.51	2.42	4.14	4.65	3.14	*	*	-
	HENBA1005755	6.51	5.76	10.77	4.34	2.96	3.2	3.26	2.6	1.41	*	*	-
	HENBA1005760	8.14	6.09	21.23	11.82	15.12	18.05	5.99	3.99	9.09			
40	HENBA1005765	16.17	18.36	30.68	12.16	17.74	19.5	18.91	12.07	18.52			
	HENBA1005766	4.59	6.7	11.4	6.4	6.48	7.14	3.63	2.36	6.46			
	HENBA1005780	66.74	88.34	50.49	70.57	83.73	79.9	51.01	23.87	50.17			
	HENBA1005795	12.49	13.53	22.26	21.04	27.08	17.3	17.89	8.53	16.93			
	HENBA1005809	27.58	28.57	37.75	10.82	11.75	10.76	7.41	20.02	13.83	**	*	-
	HENBA1005813	11.01	13.12	21.07	10.05	5.62	7.48	3.75	4.75	2.72	*	*	-
45	HENBA1005815	13.88	21.46	26.3	18.7	15.38	15.54	15.88	14.09	23.04			
	HENBA1005822	2.32	4.46	5.72	1.33	1.78	1.99	1.83	4.15	2.65			
	HENBA1005829	11.63	17.29	26.15	14.23	14.49	18.69	18.12	8.68	16.12			
	HENBA1005833	7.51	11.08	19.55	8.36	9.46	9.74	7.13	4.47	9.24			
	HENBA1005834	2.53	4.28	9.15	2.39	5.2	3.43	3.05	1.57	2.22			
	HENBA1005844	5.49	8.18	12.47	6.96	9.74	5.87	6.18	2.19	5.4			
50	HENBA1005852	9.05	12.52	17.23	5.22	7.99	5.59	7.73	5.09	7.72			
	HENBA1005853	12.09	27.26	20.62	6.51	8.22	6.09	4.4	8.87	5.67	*	*	-
	HENBA1005878	16.48	26.29	29.76	6.78	14.33	12.64	11.87	13.79	15.64	*	*	-
	HENBA1005883	10.6	8.45	21.85	5.18	5.86	5.16	3.04	2.55	2.52			
	HENBA1005884	18.4	17.45	38.78	13.72	28.17	26.47	16.14	9.17	16.98			
	HENBA1005891	14.35	23.5	29.7	12.96	17.06	15.29	15.47	12.21	17.4			
55	HENBA1005894	20.76	25	45.44	20.3	23.59	18.03	21.15	9.72	19.92			
	HENBA1005898	6.21	7.33	9.72	7.72	7.68	5.64	3.86	1.59	3.36	*	*	-
	HENBA1005902	52.63	54.68	95.93	140.97	103.88	141.36	40.77	40.69	57.15	*	+	-

Table 374

	HENBA1005313	33.27	32.37	38.89	32.29	19.06	24.67	17.8	16.01	18.82	**	-
	HENBA1005314	24.58	33.06	42.65	21.87	11.43	15.5	9.92	9.33	8.85	*	-
5	HENBA1005315	12.58	12.96	19.94	17.38	10.31	13.36	9.94	6.48	5.8	*	-
	HENBA1005317	4.9	4.68	7.03	6.55	5.75	6.41	3.83	2.01	4.66	*	-
	HENBA1005318	2.91	3.45	4.86	4.07	3.07	4.69	2.08	0.38	2.27		-
	HENBA1005324	11.54	12.46	27.19	15.78	17.61	18.74	11.73	1.86	14.23		-
	HENBA1005331	43.61	41.81	60.5	40.68	48.26	41.6	45.76	21.6	43.46		-
10	HENBA1005337	20.3	20.66	34.16	18.47	13.27	15.91	13.2	7.95	13.16	*	-
	HENBA1005338	37.76	51.08	62.83	38.86	14.3	16.63	9.18	11.59	18.35	**	-
	HENBA1005344	58.27	53.4	127.13	158.88	86.48	122.88	43.73	24.53	36.87		-
	HENBA1005353	291.67	197.22	329.41	381.89	358.33	324.87	364.08	232.83	294.3		-
	HENBA1005359	3.54	4.47	8.65	4.81	7.38	8.25	4.29	2.88	3.92		-
	HENBA1005362	14.29	12.72	20.99	11.94	11.74	13.92	13.73	6.82	10.98		-
15	HENBA1005364	3.08	8.57	11.19	9.32	9.41	5.54	6.78	3.23	3.82		-
	HENBA1005367	9.52	14.36	9.74	11.62	16.31	12.41	12.24	9.27	3.86		-
	HENBA1005372	309.34	404.59	430.49	84.13	123.69	78.11	25.56	68.19	97.12	**	-
	HENBA1005374	36.56	51.63	69.8	44.44	39.19	38.53	36.51	36.03	34.64		-
	HENBA1005379	33.5	27	43.36	22.13	30.25	17.05	21.18	29.05	27.22		-
	HENBA1005382	15.49	19.66	32.8	16.35	15.89	24.51	20.76	15.49	21.41		-
20	HENBA1005384	6.36	8.47	15.31	11.14	12.78	12.49	8.42	6.41	12.29		-
	HENBA1005386	7.43	7.3	12.47	6.27	5.44	7.51	3.39	1.08	4.63	*	-
	HENBA1005389	18.61	20.29	18.71	13.5	21.69	22.09	19.9	13.97	8.2		-
	HENBA1005394	12.89	8.67	16.27	12.3	14.5	11.32	10.11	3.58	3.69		-
	HENBA1005403	4.34	4.47	10.77	7.65	8.37	7.4	6.03	4.21	5.27		-
	HENBA1005408	5.14	5.28	8.12	4.84	5.38	6.7	5.4	4.72	1.33		-
25	HENBA1005410	147.8	198.98	244.61	96.9	175.54	169.68	68.34	63.45	81.73	*	-
	HENBA1005411	3.94	4.86	10.62	4.18	2.45	4.02	1.07	3.25	2.19		-
	HENBA1005423	7.19	6.73	19.41	6.38	7.84	8.58	6.14	3.21	5.83		-
	HENBA1005426	95.36	92.66	185.58	213.51	144.79	175.18	76.93	45.72	96.01		-
	HENBA1005427	3.18	3.68	9.55	2.9	4.08	4.42	2.98	1.86	3.96		-
	HENBA1005430	38.92	50.77	66.51	41.28	49.27	33.43	51.65	28.88	24.29		-
30	HENBA1005438	85.73	96.52	156.38	59.22	45.51	57.38	12.76	14.31	32.68	*	-
	HENBA1005443	15.2	22.18	31.05	12.38	21.42	15.9	15.18	10.34	15.89		-
	HENBA1005447	7.4	8.91	13.83	6.04	6.83	6.51	4.71	3.18	4.95	*	-
	HENBA1005449	188.67	185.78	227.88	315.2	205.18	189.94	184.99	92.08	155.69		-
	HENBA1005452	87.1	82.51	97.54	153.23	76.5	106.72	73.14	41.42	77.58		-
35	HENBA1005454	4.07	5.62	12.12	2.12	9.94	2.42	3.66	2.56	2.8		-
	HENBA1005468	3.12	5.06	12.85	2.9	5.78	4.72	3.8	1.69	1.54		-
	HENBA1005469	4.06	5.82	14.74	4.56	9.66	7.3	4.34	3.61	3.19		-
	HENBA1005472	48.84	68.93	98.37	43.92	45.45	42.16	48.44	44.29	63.31		-
	HENBA1005474	3.81	5.4	6.23	3.61	6.81	3.03	3.23	2.83	4.47		-
	HENBA1005475	18.35	13.18	29.74	13.69	11.52	8.61	14	8.86	16.92		-
	HENBA1005489	11.24	6.84	19.41	11.32	8.2	10.15	7.69	4.76	6.72		-
40	HENBA1005497	3.55	3.84	13.2	3.87	5.02	4.47	2.68	1.54	6.08		-
	HENBA1005500	9.06	12.27	15.75	5.67	8.54	10.09	5.5	7.58	6.57	*	-
	HENBA1005506	66.72	77.27	99.37	69.96	83.48	86.85	48.45	36.29	43.04	*	-
	HENBA1005508	10.52	16.74	23.57	15.52	16.57	19.17	24.35	7.4	7.32		-
	HENBA1005511	106.43	145.35	110.4	50.22	49.21	46.23	5.8	10.19	8.56	**	-
	HENBA1005513	13.22	13.61	12.43	5.61	8.6	9.42	6.36	6.5	6.97	*	-
45	HENBA1005517	8.83	13.8	21.17	12.19	9.09	9.65	12.06	10.72	8.14		-
	HENBA1005518	6.15	6.12	10.29	3.31	2.65	3.34	2.82	2.51	2.48	*	-
	HENBA1005520	4.72	5.27	15.66	1.8	9.96	5.07	11.44	4.1	6.1		-
	HENBA1005522	7.43	6.22	15.1	2.69	4.51	4.56	4.2	1.89	5.13		-
	HENBA1005526	12.95	11.98	23.79	8.97	14.67	6.23	7.41	3.51	8.41		-
	HENBA1005528	20.88	31.78	47.82	17.24	26.35	20.12	19.44	10.67	18.86		-
50	HENBA1005530	155.03	123.7	313.84	376.32	270.85	265.42	127.01	98.49	126.67		-
	HENBA1005538	4.22	2.81	4.67	3.91	2.23	1.75	0.82	1.49	1.34	*	-
	HENBA1005539	7.37	8.42	11.33	9.56	8.07	5.43	6.22	7.08	3.71		-
	HENBA1005545	6.67	6.11	8.83	3.2	2.97	2.68	4.51	2.07	2.36	*	-
	HENBA1005548	9.67	16.77	17.36	7.24	13.44	14.5	10.18	7.2	5.13		-
	HENBA1005552	51.47	60.33	76.88	44.85	67.33	45.49	61.05	38.5	41.19		-
55	HENBA1005558	1303.4	1695.5	2079.7	1863.4	2100.7	1420	1688.2	1099.2	644.47		-
	HENBA1005568	18.05	24.94	43.8	19.93	30.36	19.35	24.29	10.08	5.54		-
	HENBA1005570	9.18	10.68	10.58	5.69	4.61	5.33	3.91	1.39	3.62	**	-

Table 373

	HEMBA1004995	13.04	8.58	10.13	7.12	8.06	6.92	2.73	3.68	5.31	*	-
	HEMBA1005004	4.77	5.36	6.34	4.11	3.26	3.94	3.94	3.84	5.52	*	-
5	HEMBA1005008	57.09	40.78	98.86	67.53	69.03	54.32	23.5	39.71	21.46		
	HEMBA1005009	1057.6	723.08	1508	886.08	1346.5	1172.9	1256.5	763.02	961.92		
	HEMBA1005019	4.41	3.37	12.38	2.48	5.05	3.17	5.8	3.17	5.15		
	HEMBA1005021	20.78	39.68	34.72	6.67	27.86	14.2	11.45	16.23	4.97	*	-
	HEMBA1005029	70.74	90.66	114.47	109.76	103.35	90.04	59.53	40.06	53.52	*	-
	HEMBA1005035	18.47	18.85	35.14	26.8	19.92	22.3	9.64	15.48	14.58		
10	HEMBA1005036	6.46	8.45	12.94	4.31	7.1	4.1	3.99	2.47	4.91		
	HEMBA1005039	30.97	34.92	39.63	34.24	27.86	28.36	12.93	8.11	16.92	**	-
	HEMBA1005047	8.28	10.14	13.81	4.91	5.09	5.36	2.77	2.57	2.31	*	**
	HEMBA1005050	3.68	2.17	10.81	2.45	2.33	1.31	2.32	1.28	0.56		
	HEMBA1005062	4.16	12.91	14.13	4.05	8.34	22.08	6.52	3.12	5.74		
	HEMBA1005066	10.67	14.99	36.8	11.75	15.95	10.56	12.8	8.39	12.44		
15	HEMBA1005067	4.44	5.11	10.87	2.4	4.78	3.11	2.95	0.22	2.1		
	HEMBA1005070	52.72	54.64	132.49	164.51	146.19	162.23	48.46	33.29	55.3	*	+
	HEMBA1005075	3.89	4.62	7.08	4.3	4.86	3.7	4.12	2.52	4.18		
	HEMBA1005078	9.47	16.56	13.83	7.13	9.41	6.63	4.44	6.27	3.76	*	-
	HEMBA1005079	31.02	35.09	45.38	24.46	22.17	19.63	32.37	21.21	24.67	*	-
20	HEMBA1005083	6.87	5.26	12.68	2.37	3.44	3.16	3.82	1.83	4.73		
	HEMBA1005084	14.72	21.03	25.93	7.18	18.88	15.76	10.99	5.75	6.49	*	-
	HEMBA1005088	4.46	4.67	12.93	1.85	3.81	2.49	3.1	1.87	2.53		
	HEMBA1005089	4.71	9.88	14.88	4.89	6.06	3.51	4.1	0.92	1.92		
	HEMBA1005090	25.03	17.55	31.81	18.25	14.13	11.76	5.18	6.32	11.92	*	-
	HEMBA1005096	5.89	3.91	4.65	4.65	2.95	3.34	2.39	2.2	2.94	*	-
25	HEMBA1005101	39.08	41.35	50.86	29.48	23.15	20.02	33.26	32.78	29.58	*	*
	HEMBA1005107	40	41.43	50.54	29.82	24.91	24.18	41.74	30.86	33.12	**	-
	HEMBA1005113	113.11	107.38	156.04	92.91	86.12	76.09	106.26	47.64	78.36		
	HEMBA1005123	10.25	18.19	27.19	10.94	11.91	8.06	14	8.52	9.01		
	HEMBA1005133	17.82	23.43	41.02	14.36	24.16	22.06	27.83	16.74	25.23		
	HEMBA1005135	5.59	8.93	15.2	4.63	7.32	4.29	4.52	1.48	2.68		
30	HEMBA1005145	4.53	2.71	8.45	5.36	4.35	4.23	3.05	2.29	4.35		
	HEMBA1005149	378.91	317.95	317.16	22.18	181	208.94	102.53	216.46	124.98	*	**
	HEMBA1005152	6.82	9.2	10.54	4.39	6.2	1.79	3.59	2.97	2.92	*	**
	HEMBA1005159	7.43	8	10.73	4.97	6.07	2.08	4.16	3.29	1.57	*	-
	HEMBA1005172	7.45	10.34	13.12	4.47	4.62	4.49	4.3	1.57	2.99	*	*
35	HEMBA1005185	22.87	28.13	43.59	20.68	26.1	16.31	27.6	18.57	18.78		
	HEMBA1005186	8.26	12.01	22.97	7.71	9.71	6.32	6.15	2.19	5.11		
	HEMBA1005195	21.01	45.46	54.34	19.83	21.77	13.25	11.91	3.42	10.22	*	-
	HEMBA1005201	30.23	25.81	53.48	38.66	31.34	27.12	18.26	15.51	22.99		
	HEMBA1005202	5.59	7.98	9.58	3.99	3.6	4.2	6.81	2.2	7.96	*	-
	HEMBA1005204	16.46	12.78	28.18	11.48	14.05	9.97	14.37	10.45	14.76		
	HEMBA1005206	23.26	6.89	27.27	4.57	14.1	5.14	8.86	8.11	3.99		
40	HEMBA1005219	46.02	33.86	64.57	13.22	55.15	17.76	33.33	34.4	11.24		
	HEMBA1005223	6.35	5.89	13.54	0.44	2.75	4.32	3.73	1.82	1.78		
	HEMBA1005229	6.79	9.67	19.26	5.01	5.11	5.18	4.75	0.62	2.59		
	HEMBA1005230	211.84	132.19	314.59	37.27	92.19	87.58	104.99	40.98	12.2	*	-
	HEMBA1005232	5.02	3.23	8	4.94	3.86	4.85	3.42	2.21	2.54		
	HEMBA1005238	4.01	4.87	12.04	12.05	4.27	6.06	2.73	1.86	3.31		
45	HEMBA1005241	4.78	4.02	12.24	4.12	3.03	3.52	2.77	1.9	3.08		
	HEMBA1005244	7.96	6.13	14.8	5.66	6.64	4.51	3.63	3.58	2.47		
	HEMBA1005246	144.3	160.26	353.25	249	171.68	181.26	165.76	91.77	157.68		
	HEMBA1005251	11.91	8.98	32.78	5.64	10.3	10.41	18.87	11.34	4.1		
	HEMBA1005252	12.15	18.12	37.6	17.48	16.35	13.11	14.26	8.12	11.62		
	HEMBA1005267	41.28	35.46	59.97	65.43	48.12	53.65	34.92	15.48	29.09		
50	HEMBA1005274	146.06	122.53	200.33	113.76	78.58	123.13	106.12	96.92	123.47		
	HEMBA1005275	50.96	52.4	71.8	62.52	44.67	45.43	64.52	35.24	52.21		
	HEMBA1005288	36.19	33.62	70.75	88.7	68.37	56.26	25.84	20.1	22.79		
	HEMBA1005293	11.95	9.22	34.15	19.57	13.03	24.04	13.57	18.82	7.5		
	HEMBA1005296	17.02	8.96	24.16	11.91	18.13	12.63	5.55	16.37	7.9		
	HEMBA1005301	7.64	4.4	12.18	6.44	7.08	7.49	6.66	3.04	4.73		
55	HEMBA1005304	3.18	4.74	9.81	5.51	5.89	5.14	6.35	2.58	2.92		
	HEMBA1005305	4.23	5.99	12.03	5.04	5.33	5.38	6.54	0.53	2.54		
	HEMBA1005311	11.53	9.17	14.37	12.69	6.17	11	10.14	6.62	9.41		

Table 372

	HEMBA1004725	8.16	9.51	14.17	11.97	14.28	12.46	9.22	7.11	10.71				
	HEMBA1004730	9.5	30.77	15.01	11.4	24.39	8.96	5.69	6.95	7.79				
5	HEMBA1004733	4.35	1.94	6.11	2.44	2.1	1.31	3.15	0.66	3.23				
	HEMBA1004734	6.35	4.56	8.84	3.47	5.46	3.65	7.89	2.85	5.88				
	HEMBA1004736	16.7	24.9	39.79	8.08	17.63	7.89	15.58	8.45	13.06				
	HEMBA1004748	7.15	9.02	16.19	2.15	7.19	3.48	6.72	4.3	8.2				
	HEMBA1004749	24.36	23.03	33.63	4.23	9.92	4.61	5.83	2.01	3.6	**	**	-	-
	HEMBA1004751	2.98	5.64	15.55	3.81	7.93	4.03	5.96	2.05	2.94				
10	HEMBA1004752	114.05	106.99	178.16	68.94	72.11	86.46	30.58	32.81	51.61		*	-	-
	HEMBA1004753	63.77	72.42	98.61	28.89	47.7	46.76	41.95	35	54.38	*	*	-	-
	HEMBA1004755	14.23	13.71	21.59	11.51	11.62	10.29	10.53	11.28	11.46				
	HEMBA1004756	8	8.06	13.16	9.58	10.37	12.14	11.47	5.24	11.51				
	HEMBA1004758	28.52	33.28	46.67	20.49	28.34	19.64	31.58	21.06	23.78				
	HEMBA1004763	5.65	6.08	11.46	2.37	10.09	17.67	15.26	61.35	5.43				
15	HEMBA1004768	3.64	9.18	17.67	4.3	5.33	4.44	3.13	1.14	2.86				
	HEMBA1004770	3.52	3.97	15.56	4.59	3.64	3.31	3.4	1.61	4.49				
	HEMBA1004771	9.14	8.57	13.33	7.83	4.73	6.17	4.13	3.12	6.19	*		-	-
	HEMBA1004775	4.31	4.72	7.01	2.54	3.11	3.68	4.57	1.14	2.99				
	HEMBA1004776	24.15	24.27	34.07	11	13.72	16.67	11.79	13.87	5.14	*	*	-	-
	HEMBA1004778	7.73	3.98	15.13	3.29	12.02	3.24	6.23	3.84	2.81				
20	HEMBA1004784	24.44	36.5	43.33	23.54	19.27	15.1	33.56	28.4	19.49				
	HEMBA1004785	13.67	18.37	27.67	13.73	15.71	13.71	15.09	7.19	15.39				
	HEMBA1004789	4.14	8.07	13.1	2.66	4.24	5.12	10.75	3.53	4.53				
	HEMBA1004795	7.33	8.35	20.46	14.48	8.52	7.53	6.22	3.52	6.52				
	HEMBA1004797	7.1	4.75	10.25	8.19	4.85	5.24	4.01	3.13	5.6				
25	HEMBA1004803	8.6	9.94	11.36	7.93	6.31	7.52	6.11	2.35	5.54	*	*	-	-
	HEMBA1004806	3.19	2.79	8.69	6.88	5.26	4.11	3.33	0.92	0.9				
	HEMBA1004807	4.03	3.35	11.54	5.63	6	7.16	2.01	2.6	2.3				
	HEMBA1004816	15.48	10.97	19.39	6.27	7.54	6.26	5.63	8.15	5.23	*	*	-	-
	HEMBA1004820	7.87	10.12	20.63	12.23	11.52	10.1	9.12	3.16	8.37				
	HEMBA1004833	4.21	6.73	18.25	5.9	8.04	9.77	5.85	1.98	3.49				
30	HEMBA1004847	3.02	6.46	11.24	4.65	7.78	8.66	8.67	2.5	5.11				
	HEMBA1004850	4.4	5.74	10.78	6.32	4.58	7.77	3.54	4.67	2.65				
	HEMBA1004863	17.31	15.86	32.11	21.23	15.33	16.21	14.88	9.41	17.84				
	HEMBA1004864	17.46	20.24	32.03	15.36	13.18	16.06	19.67	7.45	13.87				
	HEMBA1004865	113.3	61.73	148.7	148.79	88.87	115.94	64.71	39.42	63.09				
	HEMBA1004880	8.83	11.17	12.28	9.73	8.92	10.05	5.47	2.68	3.9	**		-	-
35	HEMBA1004882	141.61	124.12	179	131.9	123.22	118.49	123.09	86.3	152.75				
	HEMBA1004885	7.94	7.93	12.26	7.41	11.08	9.32	5.37	3.13	5.59	*		-	-
	HEMBA1004889	35.77	44.93	60.57	51.03	25.98	30.7	17.41	45.83	36.11				
	HEMBA1004900	52.96	49.82	61.95	21.61	27.4	15.4	7.06	13.28	12.57	**	**	-	-
	HEMBA1004909	14.18	13.16	28.88	16.89	11.25	13.43	12.38	9.23	13.02				
	HEMBA1004918	6.03	4.11	8.13	4.43	3.46	3.67	3.79	1.63	1.47				
40	HEMBA1004923	9.64	9.9	14.09	8.35	13.9	10.52	5.58	4.15	7.26	*		-	-
	HEMBA1004929	3.15	3.03	7.44	3.03	3.95	3.22	2.76	0.8	3.27				
	HEMBA1004930	4.7	7.7	9.9	4.33	7.94	6.01	5.83	3.66	7.33				
	HEMBA1004933	48.9	58.84	65.65	52.38	60.44	49.38	52.85	24.83	54.95				
	HEMBA1004934	3.85	4.55	8.24	5.31	3.47	5.34	6.35	6.72	5.35				
	HEMBA1004937	12.55	13.42	25.29	16.02	10.93	15.03	11.34	9.36	13.16				
45	HEMBA1004943	6.33	9.2	8.01	2.75	3.28	2.19	4.11	2.61	3.13	**	**	-	-
	HEMBA1004944	19.58	19.31	31.82	17.17	17.7	18.72	15.62	10.66	19.2				
	HEMBA1004946	12.19	9.84	23.87	11.5	9.34	11.54	7.67	6.35	7.52				
	HEMBA1004952	5.52	5.6	7.61	2.28	4.07	4.82	1.95	1.52	2.27	**		-	-
	HEMBA1004954	95.41	161.15	61.32	125.56	130.49	124.27	79.29	36.2	131.05				
	HEMBA1004958	3.34	5.15	10.1	3.76	7.59	6.47	4.82	2.29	4.38				
50	HEMBA1004960	18.1	21	43	23.94	15.38	19.28	12.99	14.78	18.5				
	HEMBA1004971	17.31	26.71	28.24	5.08	6.02	5.94	4.63	4.75	6.7	**	**	-	-
	HEMBA1004972	122.03	238.04	172.22	44.53	137.98	72.83	61.38	80.13	88.64	*		-	-
	HEMBA1004973	3.44	2.83	7.72	3.58	1.82	6.46	2.6	2.06	3.24				
	HEMBA1004977	14	10.09	24.08	10.99	9.73	9.85	6.91	5.11	5.98				
	HEMBA1004978	11.81	9.75	18.48	9.66	9.26	8.27	8.68	4.77	8.65				
55	HEMBA1004980	6.76	8.63	15.4	3.58	6.32	6.19	5.45	3.4	4.05				
	HEMBA1004982	6.24	8	15.9	4.68	8.72	7.61	5.71	2.32	4.42				
	HEMBA1004983	27.66	31.4	50.17	31.75	22.28	27.83	26.75	25.03	37.28				

Table 371

	HEMBA1004446	1.03	3.03	4.89	2.99	6.41	1.96	3.37	0.17	1.81			
	HEMBA1004451	47.91	49.66	86.71	31.61	23.94	40.91	29.4	25.22	31.73			
5	HEMBA1004452	7.95	9.5	12.93	8.01	5.07	6.5	4.65	4.16	4.59	*	-	
	HEMBA1004454	21.61	21.48	36.91	22.76	21.2	22.64	20.66	13.58	20.72			
	HEMBA1004460	3.23	5	9.05	6.23	2.51	4.45	3.96	1.85	1.76			
	HEMBA1004461	9.82	11.87	18.26	13.46	12.46	12.25	11.75	5.13	11.97			
	HEMBA1004468	5.97	8.24	11.49	13.09	9.63	6.4	4.93	3.14	5.27			
	HEMBA1004479	7.22	6.26	11.32	4.91	8.01	9.21	5.84	3.77	4.35			
10	HEMBA1004482	5.37	8.99	11.74	7.23	3.06	5.24	4.77	2.99	3.49			
	HEMBA1004491	7.67	10.41	13.9	11.84	11.35	7.74	9.41	6.84	7.99			
	HEMBA1004499	7.91	12.25	21.91	14.4	10.84	11.41	12.99	9.9	12.88			
	HEMBA1004502	28.69	31.67	44.53	24.39	21.51	21.75	15.45	9.07	14.65	*	-	
	HEMBA1004505	9.98	9.29	16.72	11.78	9.95	10.23	12.78	4.69	8.35			
	HEMBA1004506	10.31	10.67	16.69	10.45	12.56	11.51	11.7	4.59	11.13			
15	HEMBA1004507	21.1	25.48	30.3	21.45	19.73	18.7	29.74	17.59	32.44			
	HEMBA1004509	2.41	4.16	3.19	4.33	6.97	5.13	3.86	1.09	3.24			
	HEMBA1004523	16.14	17.44	35.11	23.55	14.39	18.02	17.09	11.42	16.4			
	HEMBA1004528	10.51	13.24	19.75	9.57	16.03	13.29	6.5	10.38	8.52			
	HEMBA1004534	15.22	16.88	30.37	16.94	14.03	18.54	14.2	7.84	17.61			
20	HEMBA1004536	6.18	6.2	10.88	7.97	5.18	6.58	4.33	4.17	3.26			
	HEMBA1004538	9.62	8.85	16.21	10.28	7.48	8.38	4.46	3.25	4.87	*	-	
	HEMBA1004542	7.28	10.06	13.56	7.76	7.83	8.12	4.06	4.75	6.69			
	HEMBA1004552	3.25	4.65	8.61	4.79	5	6.2	4.33	0.53	3.64			
	HEMBA1004554	8.19	11.95	16.92	9.57	13.98	10.14	9	4.19	10.81			
	HEMBA1004558	12.36	17.94	21.39	13.09	11.7	13.96	9.64	9.22	10.83			
25	HEMBA1004560	4.66	8.68	10.86	4.85	5.89	4.95	2.86	5.27	5.83			
	HEMBA1004564	9.73	17.48	27.36	16.06	14.28	12.2	13.33	10.15	10.66			
	HEMBA1004566	3.56	3.98	12.25	2.36	4.04	2.92	3.05	2.38	2.63			
	HEMBA1004573	4.48	3.98	12.03	4.58	4.29	4.64	2.63	3.89	4.83			
	HEMBA1004576	18.09	17.62	32.93	24.33	34.17	25.83	17	11.99	13.66			
	HEMBA1004577	3.54	7.83	16.24	7.08	11.02	14.75	9.75	3.87	4.69			
30	HEMBA1004586	4.65	5.81	14.36	5.07	7.86	5.89	6.37	2	4.74			
	HEMBA1004596	284.44	249.12	425.92	413.25	261.05	315.49	139.33	269.7	335.01			
	HEMBA1004604	11.58	13.47	19.78	11.02	11.29	12.05	9.52	8.82	11.26			
	HEMBA1004607	44.74	50.81	86.88	51.55	56.7	51.72	48.7	32.51	56.83			
	HEMBA1004610	6.05	7.49	14.69	9.25	6.9	7.29	6.72	4.7	9.16			
	HEMBA1004617	12.1	9.56	20.48	8.02	10.16	8.41	9.52	5.32	9.98			
35	HEMBA1004622	5.16	6.28	10.24	4.13	6.32	4.27	7.06	2.49	4.07			
	HEMBA1004626	5.67	7.39	15	4.2	9.2	4.95	4.59	5.59	4.31			
	HEMBA1004629	13.02	21.22	41.46	19.54	21.07	18.99	20.74	13.12	14.82			
	HEMBA1004631	73.82	82.05	86.06	51.43	26.56	15.5	10.72	28.9	42.6	*	**	-
	HEMBA1004632	24.68	42.63	52.95	37.34	39.22	29.39	26.96	22.05	34.67			
	HEMBA1004633	38.98	38.11	53.52	17.88	40.68	39.12	15.84	33.55	35.48			
40	HEMBA1004636	16.64	14.75	24.41	11.5	12.22	12.63	6.1	4.97	7.02	*	-	
	HEMBA1004637	10.33	12.11	26.18	5.02	9.28	11.39	12.44	6.08	12.85			
	HEMBA1004638	17.07	21.26	52.09	19.97	27.34	24.5	24.76	14.24	22.8			
	HEMBA1004645	41.39	65.54	74.75	36.73	58.32	57.87	35.72	21.96	40.06			
	HEMBA1004656	33.89	50.4	80.19	35.73	46.51	37.77	45.77	32.71	37.95			
	HEMBA1004657	8.65	8.21	14.8	8.65	5.01	4.76	2.7	2.34	16.46			
45	HEMBA1004666	18.27	17.29	28.67	14.15	16.19	11.73	13.41	11.89	17.32			
	HEMBA1004668	9.31	10.17	13.58	7.96	6.32	5.8	4.8	3.37	4.51	*	**	-
	HEMBA1004670	6.28	8.23	12.51	5.31	4.65	7.13	5.07	1.93	2.73	*	-	
	HEMBA1004672	7.49	5.42	13.8	6.32	5.91	7.14	7.17	3.42	6.19			
	HEMBA1004689	18.06	37.4	36.02	13.85	14.84	13.32	10.59	5.45	12.39	*	-	
	HEMBA1004690	8.71	10.59	14.84	3.74	5.26	4.9	4.08	1.25	4.02	*	*	-
50	HEMBA1004693	54.63	51.62	109.86	81.26	86.47	81.99	74.06	35.68	58.9			
	HEMBA1004697	107.99	90.54	174.23	129.1	114.25	71.95	45.98	67.21	100.42			
	HEMBA1004702	17.82	20.4	33.62	24.43	14.82	19.64	15.1	13.79	12.53			
	HEMBA1004704	4.12	4.28	3.95	3.01	2.64	2.5	2.6	1.84	1.67	**	**	-
	HEMBA1004705	8.53	8.58	10.66	4.87	5.69	4.53	7.25	3.35	5.23	**	*	-
	HEMBA1004706	4.44	4.15	10.51	3.36	3.58	2.66	4.3	1.58	2.61			
55	HEMBA1004709	17.24	19.06	37.38	23.46	18.59	23.59	17.98	10.66	14.19			
	HEMBA1004711	162.08	157.26	293.47	260.03	204.86	251.04	138.17	104.82	96.04			
	HEMBA1004723	8.81	12.37	23.58	15.07	18	14.65	13.5	4.03	7.35			

Table 370

	HEMBA1004207	5.23	7	7.18	5.06	6.93	4.27	6.27	2.78	3.75				
	HEMBA1004210	4.96	5.53	8.99	5.1	8.24	7.88	8.78	4.25	6.82				
5	HEMBA1004225	11.76	16.11	15.08	1.96	9.21	5.85	1.74	2.95	4.57	*	**	-	-
	HEMBA1004227	5.13	5	14.56	6.73	5.83	3.8	3.56	2.92	3.47				
	HEMBA1004235	55.87	67.36	100.86	60.82	63.74	64.98	56.93	44.28	61.25				
	HEMBA1004237	63.06	67.44	67.86	54.44	56.68	54.92	61.37	42.55	49.6	**		-	-
	HEMBA1004238	5.4	9.74	19.37	3.19	7.79	5.08	4.55	3.62	5.8		*	-	-
10	HEMBA1004241	58.53	94.19	86.84	45.42	61.25	30.71	41.3	31.18	35.09				
	HEMBA1004242	95.96	94.64	132.52	129.29	124.32	88.47	86.26	40.26	43.03				
	HEMBA1004243	7.18	9.59	11.11	5.01	7.55	7.79	4.07	6.19	8.9				
	HEMBA1004246	3.92	5.45	6.92	3.64	1.73	7.61	2.23	1.7	3.01	*	**	-	-
	HEMBA1004247	36.08	39.65	23.46	13.27	21.43	17.87	8.15	9.3	7.87	*	**	-	-
	HEMBA1004248	60.7	102.92	68.75	40.08	49.28	54.01	55.12	49.8	49.38				
	HEMBA1004250	6.46	7.1	17.92	5.92	8.57	3.92	3.52	3.06	5.66	*		-	-
15	HEMBA1004252	9.76	13.49	18.16	5.3	8	5.39	8.18	4.43	8.93	*		-	-
	HEMBA1004260	3.71	7.97	10.5	2.84	9.75	1.75	6.82	1.55	1.83		*	-	-
	HEMBA1004264	4.86	4.92	5.04	3.34	5.85	4	4.03	2.37	1.97		*	-	-
	HEMBA1004267	14.31	26.05	30.36	19.35	16.25	13.91	15.22	15.98	20.33	**	**	-	-
	HEMBA1004272	7.23	8.73	9.64	3.95	3.4	3.74	3.02	1.81	2.67				
20	HEMBA1004274	3.8	4.23	13.32	4.59	3.82	2.88	5.15	2.11	5.65				
	HEMBA1004275	6.92	5.28	11.01	3.44	3.27	3.1	4.62	3.75	4.56		*	-	-
	HEMBA1004276	53.98	104.51	88.12	40.54	57.2	47.57	40.16	23.52	29.44				
	HEMBA1004279	3.9	4.4	11.72	1.47	4.01	1.18	2.34	1.51	2.01				
	HEMBA1004284	15.6	17.68	32.1	11.85	15.45	12.29	19.44	11.68	15.92				
	HEMBA1004286	1.1	4.52	5.35	1.91	3.11	0	3.7	0.1	2.34				
25	HEMBA1004289	4.33	10.31	14.89	4.93	9.21	6.85	5.43	4.43	9.27				
	HEMBA1004293	14.22	23.33	31.53	19.23	14.77	12.78	16.77	15.46	15.6				
	HEMBA1004295	5.99	9.65	12.39	6.69	4.97	4.03	6.51	6.53	6.32				
	HEMBA1004302	27.72	69.78	67.78	35.38	30.81	31.86	31.74	25.34	24.05				
	HEMBA1004306	22.59	41.69	39.11	30.27	34.77	20.17	21.64	11.35	18.3				
	HEMBA1004312	8.29	12	23.23	5.76	9.79	3.74	7.73	4.58	8.66				
30	HEMBA1004314	17.03	23.79	39.64	14.71	20.95	13.02	20.7	16.59	24.72				
	HEMBA1004321	2.16	4.06	8.11	3.21	4	1.12	2.54	0	3.38				
	HEMBA1004323	4.43	3.01	0.68	0.1	10.07	1.6	2.71	1.11	0.96				
	HEMBA1004327	3.2	5.57	7.13	3.58	3.76	2.03	3.61	2.13	4.27				
	HEMBA1004329	47.09	53.32	108.52	56.12	44.95	40.88	71.49	47.58	48.41				
	HEMBA1004330	7.28	14.18	16.85	8.09	8.07	7.74	11.04	9.62	5.53				
35	HEMBA1004334	6.28	6.86	13.5	5.07	6.47	3.23	5.33	2.42	4.72				
	HEMBA1004335	2.77	3.49	11.65	1.67	3.64	0.75	1.36	0.2	2				
	HEMBA1004341	4.79	7.07	13.59	3.55	3.83	2.11	5.23	1.8	5.75				
	HEMBA1004344	5.88	15.53	22.36	10.54	9.43	7.32	16.14	4.47	6.61				
	HEMBA1004347	8.48	17.07	21.37	11.69	14.36	7.58	5.26	6.18	12.85				
	HEMBA1004349	19.03	22.2	39.78	17.07	23.84	11.42	16.44	11.72	16.17				
40	HEMBA1004352	30.35	31.05	54.47	23.2	27.22	18.91	26.94	20.35	19.47				
	HEMBA1004353	65.35	90.37	111.41	40.12	57.25	44.69	62.88	49.41	51.19	*		-	-
	HEMBA1004354	31.04	48.91	55.45	29.8	26.92	23.18	26.7	16	25	*		-	-
	HEMBA1004356	36.8	69.45	87.9	42.77	38.63	30.42	48.78	31.27	34.28				
	HEMBA1004360	21.37	39.1	60.88	24.3	25.32	19.97	35.28	18.19	17.1				
	HEMBA1004366	3.36	19.8	13.89	9.13	8.25	3.58	5.88	3.78	6.29				
45	HEMBA1004372	25.71	35.24	61.82	27.78	27.66	28.9	23.52	16.32	31.11				
	HEMBA1004377	57.43	64.54	125.12	52.77	52.36	39.76	63.76	39.09	45.41				
	HEMBA1004389	26.6	23.69	56.36	35.4	23.53	18.76	21.22	13.56	17.27				
	HEMBA1004391	15.02	20.13	38.52	17.1	15.91	14.1	20.59	16.16	16.69				
	HEMBA1004393	6.23	8.43	14.77	10.01	2.68	5.81	6.08	4.76	4.24				
	HEMBA1004394	6.59	9.68	22.73	7.13	7.36	4.87	8.86	2.4	3.32				
	HEMBA1004396	6.98	7.79	21.04	5.3	10.85	6.63	4.46	2.99	4.78				
50	HEMBA1004401	25.42	37.64	61.19	35.6	34.11	30.34	50.39	18.9	33.72				
	HEMBA1004405	12.61	20.04	37.41	22.75	14.99	14.73	15.19	9.3	13.54				
	HEMBA1004408	33.43	47.3	70.7	34.73	36.57	32.16	33.47	25.45	21.56				
	HEMBA1004414	11.75	12.41	34.08	20.66	20.73	15.48	9.96	9.26	12.21				
	HEMBA1004429	11.38	11.1	25.62	16.77	12.96	11.07	10.66	8.23	8.04				
55	HEMBA1004433	4.53	4.23	9.92	6.93	6.74	8.35	4.52	0.98	3.02				
	HEMBA1004440	12.02	23.81	31.06	19.67	29.44	19.7	15.12	10.66	15.31				
	HEMBA1004444	18.31	24.79	46.54	21.84	22.35	18.04	25.64	13.85	21.08				

Table 369

	HENBA1003959	14.89	19.53	30.66	18.5	16.34	12.43	19.48	10.78	14.94			
	HENBA1003960	6.1	5.31	11.07	8.55	9.98	6.62	4.38	4.38	3.84			
5	HENBA1003966	4.32	4.36	6.33	4.53	3.74	1.94	2.49	2.11	0.85	*	-	
	HENBA1003967	11.38	13.34	19.36	11.07	6.9	6.77	8.93	7.19	5.9	*	-	
	HENBA1003968	52.83	102.28	87.2	14.25	26.41	16.31	15.76	9.02	11.36	*	**	-
	HENBA1003974	4.37	4.4	7.25	2.02	4.53	1.13	3.82	0.89	2.56			
	HENBA1003976	15.15	20.18	27.02	12.74	18.84	11.73	18.38	14.83	18			
	HENBA1003977	8.76	9.42	22.28	9.64	11.17	3.97	8.14	5.18	9.28			
10	HENBA1003978	9.01	16.51	20.07	10.86	6.71	7.75	11.84	5.73	13.48			
	HENBA1003981	68.06	77.17	117.23	62.61	51.44	47.41	51.47	46.06	74.31			
	HENBA1003982	5.4	13.31	18.21	3.94	11.62	6.14	3.92	4.82	3.97			
	HENBA1003985	7.59	8.14	15.01	6.55	7.27	6.74	5.12	5.29	7.02			
	HENBA1003987	10.14	14.01	20.78	6.87	9.2	5.37	6.58	6.5	8.96			
	HENBA1003989	10.11	15.44	20.52	5.53	5.4	4.86	8.66	4.07	5.03	*	*	-
15	HENBA1004000	2.93	4.96	11.45	2.25	2.97	1.76	2.52	0.75	2.54			
	HENBA1004006	18.54	20.18	42.2	18.19	19.39	15.33	18.84	12.3	17.01			
	HENBA1004007	19.65	39.73	40.28	26.62	20.72	17.41	38.95	21.37	27.19			
	HENBA1004010	67.05	91.68	2.46	0.19	0.95	0.35	50.08	44.9	72.79			
	HENBA1004011	10.54	11.89	17.43	5.17	7.46	5.81	5.72	5.31	4.01	*	*	-
20	HENBA1004012	20.42	22.3	36.75	7.27	17.23	8.75	11.84	7.16	6.96	*	*	-
	HENBA1004015	4.26	3.69	8.92	3.2	3.73	2.56	1.97	1.37	3.1			
	HENBA1004024	7.24	8.25	20.14	4.09	6.96	4.29	7.49	2.93	4.36			
	HENBA1004029	3.05	5.65	11.19	0.74	1.77	1.68	2.55	0.21	0.47			
	HENBA1004038	30.88	38.16	65.02	18.09	23.34	18.57	34.1	18.52	13.77			
	HENBA1004042	0.37	2.38	5.48	0.32	1.77	1.14	0.62	0	0.22			
25	HENBA1004045	20.4	33.53	50.23	28.78	29.43	29.29	14.02	11.2	16.1			
	HENBA1004048	37.85	38.8	52.99	50.26	40.74	37.18	34.52	20.41	13.02			
	HENBA1004049	21.46	22.41	47.43	24.15	22.42	24.41	19.84	20.47	15.98			
	HENBA1004051	9.9	8.46	21.25	12.84	10.63	10.43	12.39	7.87	8.34			
	HENBA1004053	15.29	25.2	37.82	19.98	13.35	13.99	21.18	11.93	18.89			
	HENBA1004055	5.25	11.05	5.64	0.16	1.02	0	9.11	3.17	5.22	*	-	
30	HENBA1004056	2.41	5.85	10.01	3.45	3.17	2.37	4.07	0.71	2.14			
	HENBA1004060	6.43	8.34	15.71	14.49	7.75	6	14.41	5.93	6.72			
	HENBA1004061	7.43	8.52	15.21	11.54	9.65	13.29	5.83	3.71	5.22			
	HENBA1004067	8.97	11.51	14.78	16.09	9.22	11.3	7.43	5.81	9.07			
	HENBA1004071	2.92	2.4	5.93	4.14	2.31	3.09	3.1	0.97	2.2			
	HENBA1004074	126.78	177.45	131.41	229.36	160.88	89.6	187.87	113.1	134.85			
35	HENBA1004078	20.55	20.84	34.63	28.09	31.27	21.15	16.53	16.06	27.35			
	HENBA1004085	20.65	31.8	30.03	32.27	21.47	13.85	11.87	12.81	9.59	*	-	
	HENBA1004086	2.06	3.76	5.23	2.03	3.33	2.95	2.57	0.6	1.89			
	HENBA1004097	1.94	3.31	1.59	1.13	4.7	3.06	2.83	1.2	1.23			
	HENBA1004100	14.77	24.96	29.01	20.87	18.7	13.96	10.94	12.24	15.57			
	HENBA1004103	7.32	13.02	15.94	12.18	10.65	9.17	7.19	6.37	7.49			
40	HENBA1004110	19.03	26.3	37.33	22.14	20.14	20.02	18.64	15.91	19.14			
	HENBA1004111	24.42	41.97	41.29	22.57	21.56	20.91	28.72	16.68	19.6			
	HENBA1004124	10.47	10.61	15.29	11.7	10.51	11.56	7.45	6.39	9.02			
	HENBA1004130	6.32	9.6	13.12	7.07	7.66	3.98	10.02	6.52	8.94			
	HENBA1004131	11.62	15.68	22.49	12.3	15.58	10.2	7.69	8.91	12.88			
	HENBA1004132	1.7	3.66	8.07	2.75	4.82	2.5	5.8	1.43	6.28			
45	HENBA1004133	61.09	100.97	106.22	48.94	25.58	18.64	11.83	36.24	13.37	*	*	-
	HENBA1004138	8.15	8.84	9.84	8.72	6.31	5.94	4.18	4.48	5.03	**	-	
	HENBA1004143	6.39	11.32	13.28	4.24	5.74	4.33	4.63	4.09	3.58	*	-	
	HENBA1004146	12.23	19.23	24.6	16.14	12.69	10.61	14.04	14.11	16.75			
	HENBA1004148	5.02	6.85	14.92	7.36	6.63	8.78	2.34	5.06	6.01			
	HENBA1004149	13.01	27.48	21.59	12.24	21.36	25.46	6.33	6.45	12.24			
50	HENBA1004150	162.3	225.69	242.62	246.24	265.35	201.29	161.22	151.68	209.57			
	HENBA1004154	54.95	73.81	77.08	67.59	61.25	43.45	53.14	35.47	45.98	*	-	
	HENBA1004164	27.89	39.95	75.25	53.13	38	41.78	27.04	21.48	32.66			
	HENBA1004168	27.29	41.16	48.5	30.28	31.63	25.87	20.87	19.25	25.4			
	HENBA1004199	11.53	17.04	30.46	12.77	16.45	14.72	15.81	10.26	14.34			
	HENBA1004200	10.21	15.6	23.4	12.76	12.18	11.22	9.51	9.03	10.57			
55	HENBA1004201	27.51	39.39	51.71	29.35	27.61	28.68	24.64	21.71	24.17			
	HENBA1004202	18.3	30.46	37.1	17.92	20.38	16.33	20.83	15.59	21.86			
	HENBA1004203	11.35	16.48	25.18	12.27	10.37	8.08	12.69	8.41	9.37			

Table 368

	HEMBA1003679	15.99	25.43	34.72	23.29	23.03	14.07	32.67	15.26	20.68			
	HEMBA1003680	2.36	1.32	3.39	2.39	1.66	4.26	2	1.23	1.59			
5	HEMBA1003684	27.44	32.04	37.17	24.58	15.64	21.48	12.03	9.61	13.69	*	**	-
	HEMBA1003690	26.36	32.22	54.74	28.94	29.04	32.33	34.36	26.69	39.53			
	HEMBA1003692	6.49	10.11	9.34	7.04	5.15	6.04	7.51	6.23	5.84			
	HEMBA1003702	8.56	9.3	12.43	12.4	14.68	13.17	6.17	5.02	7.69			
	HEMBA1003711	8.11	10.45	15.16	11.96	9.29	6.3	7.03	5.46	7.91			
10	HEMBA1003714	6.44	11.03	12.83	2.97	13.46	9.98	7.86	7.19	9.67			
	HEMBA1003715	14.81	13.59	14.56	12.43	13.78	10.5	10.14	8.94	12.29	*		-
	HEMBA1003717	42.45	55.77	82.27	41.74	41.39	40.14	43.46	40.44	41.87			
	HEMBA1003720	6.21	6.28	13.24	9.93	12.72	5.69	6.86	5.05	6.12			
	HEMBA1003725	3.95	3.77	7.26	6.08	3.65	3.36	2.76	2.23	2.52			
	HEMBA1003728	4.28	12.26	8.88	10.99	2.88	4.38	6.4	1.69	3.67			
15	HEMBA1003729	12.13	13.18	16.13	8.92	13.67	11.13	18.08	6.56	14.8			
	HEMBA1003732	30.17	53.31	40.17	33.85	38.01	30.07	19.05	11.75	17.73	*		-
	HEMBA1003733	3.45	4.52	4.35	2.89	6.7	2.94	3.18	1.76	2.64	*		-
	HEMBA1003742	9.01	9.99	10.98	9.56	10.71	5.69	9.3	5.6	9.68			
	HEMBA1003743	9.87	14.77	20.42	19.53	13.74	11.37	9.02	7.5	9.36			
	HEMBA1003758	14.27	20.31	26.23	15.6	15.1	13.01	12.22	13.09	13.08			
20	HEMBA1003760	19.21	17.1	29.51	6.01	14.54	12.36	3.4	6.9	5.12	*		-
	HEMBA1003764	14.01	17.88	25.76	13.37	20.5	16.53	15.36	15.5	19.31			
	HEMBA1003769	38.51	54.43	53.66	30.93	34.09	32.37	42.08	31.8	46.8	*		-
	HEMBA1003773	6.19	9.28	8.37	4.69	7.62	4.92	2.95	1.6	4.58	*		-
	HEMBA1003783	2.33	2.73	7.18	6	7.98	9.31	5.24	2.04	3.66			
	HEMBA1003784	24.68	20.11	23.17	22.69	20.48	14.65	16.36	8.3	11.94	*		-
25	HEMBA1003794	35.77	59.53	74.01	32.36	32.73	29.52	23.99	30.47	35.64			
	HEMBA1003799	28.51	43.08	44.43	24.39	25.46	22.11	20.76	21.74	26.31	*	*	-
	HEMBA1003803	6.9	7.98	12.5	2.71	3.98	5.61	1.91	2.94	2.14	*		-
	HEMBA1003804	12.17	15.5	24.58	5.01	10.25	5.91	7.51	11.06	8.61			
	HEMBA1003805	15.63	21.11	29.95	13.7	15.15	11.89	15.1	12.06	15.76			
	HEMBA1003807	12.24	15.49	18.72	15.37	14.25	10.11	18.36	12.3	19.08			
30	HEMBA1003810	4.78	6.92	11.21	4.53	9.91	7.45	6.79	2.53	4.42			
	HEMBA1003827	10.51	13.97	18.38	12.97	15	8.61	18.16	7.58	9.11			
	HEMBA1003836	4.91	7.17	10.85	4.73	8.8	6.53	3.27	4.61	5.48			
	HEMBA1003838	48.74	66.87	76.01	40.91	58.53	48.34	31.6	26.6	30.51	*		-
	HEMBA1003843	23.09	39.55	55.11	24.97	26.61	27.21	27.17	12.95	30.84			
	HEMBA1003846	18.74	19.56	28.45	26.4	19.75	21.25	14.77	12.94	15.9			
35	HEMBA1003856	31.17	48.34	48.82	28.74	30.87	25.87	49.83	42.07	42.91			
	HEMBA1003857	3.66	4.84	10.52	2.38	4.43	1.85	3.66	2.4	2.28			
	HEMBA1003864	24.35	38.16	29.59	24.03	32.07	18.61	15.17	4.79	6.89	*		-
	HEMBA1003866	15.55	24.7	24.92	21.6	24.17	13.88	22.15	14.95	14.16			
	HEMBA1003868	3.42	4.24	4.55	5.4	4.56	2.52	1.88	2.32	3.07	*		-
	HEMBA1003879	54.55	71.41	99.07	52.85	60.23	59.91	38.15	32.08	50.85			
40	HEMBA1003880	26.75	40.67	50.47	32.31	30.57	23.71	28.74	17.94	31.69			
	HEMBA1003884	3.51	2.7	10.4	2.34	3.19	1.65	5.36	1.65	2.02			
	HEMBA1003885	15.83	16.07	23.65	9.67	12.61	9.91	13.22	10.04	13.72	*		-
	HEMBA1003887	8.92	10.53	15.63	3.65	12.51	2.64	7.43	4.94	3.2			
	HEMBA1003890	5.77	11.6	13.5	3.76	5.14	4.95	4.38	2.51	3.42	*		-
	HEMBA1003893	2.5	2.24	6.97	3.71	6.34	3.91	2.06	0.55	2.76			
45	HEMBA1003896	13.16	13.74	38.07	17.77	20.7	14.83	10.45	11	13.75			
	HEMBA1003902	6.09	5.71	13.87	4.58	5.68	4.39	3.52	2.56	5.08			
	HEMBA1003904	41	55.7	67.32	43.67	30.73	35.35	31.08	26.57	32.22	*		-
	HEMBA1003908	27.42	42.25	49.6	27.37	22.92	19.91	28.26	21.38	25.22			
	HEMBA1003926	35.54	56.96	89.19	56.48	59.31	44	48.79	37.72	50.83			
	HEMBA1003937	14.37	14.68	20.38	10.43	17.34	12.15	14.47	8.13	15.59			
50	HEMBA1003939	3.4	3.02	10.99	1.88	5.36	0.25	1.12	0.35	2.25			
	HEMBA1003940	6.76	11.65	10.45	4.59	10.56	5.86	9.9	2.16	3.94			
	HEMBA1003941	31.48	46.87	93.42	56.75	59.75	51.31	23.68	24.05	38.11			
	HEMBA1003942	3.19	4.4	9.44	4.64	4.07	4.72	1.95	2.11	3.24			
	HEMBA1003945	59.01	80.97	80.84	37.7	37.04	46.98	37.87	33.83	28.54	*	**	-
	HEMBA1003949	12.02	16.53	20.22	10.36	8.89	9.69	11.57	6.85	10.8			
55	HEMBA1003950	27.53	47.63	43.7	25.53	28.11	14.15	20.64	11.67	12.95	*		-
	HEMBA1003953	10.33	12.12	25.01	8.15	9.43	6.53	13.65	9.93	7.98			
	HEMBA1003958	4.65	6.88	15.41	6.73	7.93	3.72	5.9	3.43	4.74			

Table 367

	HENBA1003418	9.46	10.25	16.29	13.88	11.67	7.88	6.52	7.48	6.95		
	HENBA1003420	15.13	21.37	34.54	8.98	7.12	8.99	3.93	4.32	8.36	*	-
5	HENBA1003425	3.65	4.02	7.55	3.03	3.24	2.46	1.29	1.12	2.83		
	HENBA1003433	12.41	15.45	35.09	18.37	16.85	15.26	10.85	5.12	17.58		
	HENBA1003440	22.95	34.38	33.51	20.78	28.45	21.36	23.78	22.34	28.46		
	HENBA1003442	12.86	16.85	24.45	11.09	13.33	10.29	12.08	9.47	11.31		
	HENBA1003447	27.49	33.89	33.23	17.4	23.73	18.49	24	17.15	19.81	*	*
	HENBA1003453	4.92	6.64	13.34	4.01	5.69	5.13	2.36	4.48	3.86		
10	HENBA1003461	31.42	55.4	52.29	25.16	39.67	29.85	35.26	26.74	37.73		
	HENBA1003463	20.03	30.11	90.85	47.21	1.41	0.3	30.58	24.08	33.93		
	HENBA1003465	4.42	6.66	9.68	7.8	5.17	6.44	3.4	2.84	6.1		
	HENBA1003480	8.18	9.01	15.32	7.58	7.38	7.85	10.22	6.26	8.1		
	HENBA1003485	8.87	6.95	9.95	4.15	3.42	4.03	5.45	1.61	3.1	**	*
	HENBA1003487	40.56	64.18	72.24	34.53	33.37	41.02	45.43	44.42	46.48		
15	HENBA1003492	47.07	69.63	47.91	15.87	60.76	43.2	54.01	24.37	34.94		
	HENBA1003494	8.97	13.82	17.09	5.4	7.03	7.86	9.14	5.83	9.54		
	HENBA1003497	27.59	34.3	36.93	14.93	24.2	20.37	27.68	19.83	25.35	*	-
	HENBA1003503	17.58	24.95	1.06	0.82	1.26	0.04	6.53	10.32	21.23		
	HENBA1003511	10.13	12.6	23.58	12.03	11.2	10.45	9.65	8.73	14.36		
20	HENBA1003528	10.75	13.6	9.55	5.22	4.34	5.54	2.88	3.04	2.97	**	**
	HENBA1003530	6.02	6.59	12.41	4.68	4.85	4.75	5.24	2.55	3.96		
	HENBA1003531	12.73	22.2	30.65	9.86	14.37	9.03	13.96	8.06	18.14		
	HENBA1003532	13.65	14.91	22.65	9.4	21.22	7.2	8.43	8.83	11.91		
	HENBA1003538	10.23	12.68	27.62	8.9	13.63	10.34	9.68	5.53	8.3		
	HENBA1003545	4.35	9.29	13.49	7.12	6.86	5.1	7.44	2.65	6.35		
25	HENBA1003546	6.53	3.47	17.38	20.26	8.26	12.59	2.96	5.15	4.61		
	HENBA1003548	2.71	6.58	8.78	3.2	9.19	3.14	6.24	4.03	2.98		
	HENBA1003553	4	4.81	7.29	3.53	3.84	2.73	2.82	2.01	2.64	*	-
	HENBA1003555	8.02	14.32	16.54	7.4	9.1	7.1	7.78	7.09	7.03		
	HENBA1003556	5	8.35	8.92	2.85	4.91	3.37	2.9	2.25	2.94	*	-
	HENBA1003560	11.54	14.47	20.06	8.38	13.16	7.52	7.48	5.19	5.3	*	-
30	HENBA1003565	12.47	6.48	14.4	4.78	7.63	5.19	7.77	3.47	4.1		
	HENBA1003568	167.82	373.84	257.14	82.85	193.26	96.08	64.68	26.85	13.07	*	-
	HENBA1003569	19.02	23.29	63.64	40.15	29.03	28.36	20	16.6	26.95		
	HENBA1003571	8.12	11.43	9.83	5.26	10.52	5.78	8.95	5.27	5.01		
	HENBA1003579	22.13	20.38	39.63	17.94	15.09	12.56	17.53	15.68	17.94		
	HENBA1003580	18.63	22.77	31.1	30.35	20	21.57	12.72	8.11	11.25	*	-
35	HENBA1003581	4.54	7.44	11.51	2.94	2.83	4.01	3.4	1.33	1.8		
	HENBA1003591	9.77	30.24	25.57	6.75	16.55	5.72	11.87	5.17	9.69		
	HENBA1003595	11.6	18.67	39.48	13.28	20.4	12.94	13.52	5.6	8.35		
	HENBA1003597	34.44	63.26	65.19	37.08	35.21	30.06	23.33	8.38	16.41	*	-
	HENBA1003598	2.25	2.74	3.02	2.41	2.07	1.24	1.39	0.6	1.38	*	-
	HENBA1003600	21.91	26.66	48.26	21.72	21.86	19.48	19.9	18.88	27.46		
40	HENBA1003602	5.44	4.97	13.36	4.53	6.21	5.48	3.61	3.19	3.61		
	HENBA1003604	4.52	2.99	9.05	3.91	3.29	1.56	2.58	2.5	3.04		
	HENBA1003610	13.16	15.85	27.61	10.3	9.81	5.68	10.32	8.77	8.07		
	HENBA1003615	14.12	19.51	37.31	8.32	15.31	10.64	25.53	7.2	9.53		
	HENBA1003617	2.15	10.76	12.38	1.63	5.11	1.57	4.87	0.26	2.23		
	HENBA1003620	1.75	4.53	12.78	1.34	3.78	1.52	1.5	0.27	0.68		
45	HENBA1003621	12.05	16.03	37.29	21.96	16.06	15.37	10.88	5.36	14.36		
	HENBA1003622	7.66	5.59	10	3.27	5.47	4.78	6.97	4.55	2.99		
	HENBA1003630	43.11	47.17	86.26	32.33	46.01	44.24	29.63	21.05	20.55		
	HENBA1003637	820.66	472.86	680.3	365.52	904.5	411.7	621.11	976.32	238.58		
	HENBA1003640	62.65	69.86	125.99	45.58	48.68	42	65.31	45.48	45.21		
	HENBA1003645	15.97	22.75	34.7	19.9	17.55	14.87	21.4	9.64	18.1		
50	HENBA1003646	6.76	13.27	28.86	11.35	10.28	4.41	12.2	6.62	10.03		
	HENBA1003647	68.54	91.82	142.01	91.35	74.78	80.66	83.06	37.92	60.33		
	HENBA1003656	41.62	46.94	78.5	39.66	28.08	37.31	18.6	12.53	17.55	*	-
	HENBA1003662	9.43	10.68	20.41	19.54	13.98	13.03	9.65	4.93	11.88		
	HENBA1003666	8.44	8.9	21.02	18.11	19.64	14.87	11.75	6.74	12.93		
	HENBA1003667	27	32.49	46.33	29.7	40.13	34.43	25.82	21.77	25.87		
55	HENBA1003670	16.71	15.12	35.71	10.11	19.12	14.79	10.97	18.94	10.06		
	HENBA1003674	3.06	3.93	11.38	12.14	7.69	5.61	4.08	1.48	5.42		
	HENBA1003677	3.62	4.3	9.92	2.94	5.26	2.08	4.03	0.72	1.84		

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	HEMBA1003152	4.25	6.9	16.54	3.43	9.18	3.49	6.82	2.75	2.37			
	HEMBA1003157	16.47	20.65	36.14	14.57	15.21	12.8	18.89	12.37	10.25			
5	HEMBA1003186	6.07	8.14	15.45	9.26	7.78	7.15	6.88	3	7.14			
	HEMBA1003171	4.76	6.01	1.83	0.22	9.4	0.95	4.99	3.3	4.74			
	HEMBA1003175	9.06	9.36	15.39	5.94	5.41	4.09	8.31	5.54	6.76	*	-	
	HEMBA1003179	11.74	19.14	23.08	11.43	19.69	16.88	15.24	11.13	8.75			
	HEMBA1003186	32.34	34.21	53.13	30.02	30.6	23.62	32.72	27.67	23.79			
10	HEMBA1003196	4.93	5.75	11.29	1.52	4.36	1.61	2.88	1.66	1.27			
	HEMBA1003197	11.79	16.34	26.33	10.88	17.37	11.43	13.65	9.71	13.56			
	HEMBA1003199	23.1	49.93	40.51	25.63	28.94	20.25	30.98	8.01	23.38			
	HEMBA1003202	23.9	28.7	33.49	4.92	13.85	16.67	6.88	7.46	15.28	*	**	-
	HEMBA1003204	4.47	10.56	14.07	5.38	6.97	3.66	4.1	2.93	5.45			
	HEMBA1003210	26.93	30.85	39.03	16.12	18.97	10.78	20.75	18.2	19.07	*	*	-
15	HEMBA1003212	25.47	38.89	40.06	19.34	20.63	15.78	34.72	24.84	21.07	*	-	
	HEMBA1003218	4.32	4.85	11.19	2.63	4.58	1.07	3.74	6.58	2.3			
	HEMBA1003220	35.02	49.9	56.61	20.36	27.56	20.54	37.25	33.59	30.02	*	-	
	HEMBA1003222	6.38	11.47	27.57	6.14	9.88	5.85	9.63	5.71	6.41			
	HEMBA1003225	1.99	5.89	11.4	3.67	2.98	2	2.78	0.36	1.82			
	HEMBA1003229	4.11	5.35	10.53	5.06	6.91	5.05	3.62	3.37	4.28			
20	HEMBA1003230	19	22.89	41.86	21.05	18.18	18.78	15.9	9.68	15.54			
	HEMBA1003235	6.36	6.82	18.89	3.6	5.78	3.31	5.14	2.94	2.76			
	HEMBA1003236	28.57	38.02	51.11	22.25	24.92	17.63	25.41	23	19.94			
	HEMBA1003250	3.88	5.03	10.29	3.12	4.49	3.35	3.78	2.8	3.87			
	HEMBA1003252	10.38	11.26	18.86	6.09	9.53	8.62	8.03	4.87	3.9			
	HEMBA1003257	4.81	5.23	14.11	3.29	4.4	2.15	4.08	0.86	2.62			
25	HEMBA1003268	4.15	9.87	10.08	10.51	3.68	3.28	8.7	2.22	2.32			
	HEMBA1003273	11.8	16.03	28.72	15.07	19.11	17.4	12.91	12.48	21.18			
	HEMBA1003276	149.48	149.2	143.91	64.58	44.3	74.17	103.96	79.83	39.01	**	*	-
	HEMBA1003277	46.25	23.14	67.56	22.07	19.01	38.9	14.72	11.37	3.34			
	HEMBA1003278	4.71	4.24	12.6	4.3	4.28	2.48	2.75	2.48	1.06			
	HEMBA1003280	129.77	109.77	149.42	22.66	42.91	45.72	29.75	44.29	19.43	**	**	-
30	HEMBA1003281	66.35	33.44	113.06	20.36	31.97	38.12	40.97	35.41	13.12			
	HEMBA1003284	4.17	19.37	13.16	7.85	6.12	2.69	7.14	1.71	3.34			
	HEMBA1003286	19.63	12.62	36.72	8.93	14.12	12.7	52.4	15.77	36.62			
	HEMBA1003291	72.44	65.02	121.3	168.07	81.81	87.09	38.11	45.46	43.73			
	HEMBA1003294	2.81	2.31	6.5	6.68	2.5	2.72	2.16	1.36	1.58			
	HEMBA1003296	44.35	42.6	76.86	89.38	53.17	64.95	32.4	28.59	38.37			
35	HEMBA1003304	9.89	11.83	11.34	8.37	7.94	10.02	7.8	7.09	8.26	**	-	
	HEMBA1003306	13.02	19.68	19.63	24.93	17.19	18.48	10.47	8.08	13.83			
	HEMBA1003309	1.78	3.09	6.44	4.38	2.65	4.86	2.49	0.95	2.26			
	HEMBA1003314	2.38	3.32	2.13	4.26	4.38	3.35	2.15	1.3	2.04	*	+	-
	HEMBA1003315	4.7	9	9.09	4.13	2.95	4.64	2.23	3.29	3.07	*	*	-
	HEMBA1003322	21.51	21.29	36.71	38.62	23.41	31.81	14.65	15.44	16.23			
40	HEMBA1003326	6.02	5.69	11.31	6.59	5.79	5.51	2.11	1.05	2.65	*	-	
	HEMBA1003327	7.54	6.06	15.43	8.47	7.53	8.8	4.99	4.52	3.75			
	HEMBA1003328	105.65	106.38	147.91	165.04	127.51	111.83	74.7	76.39	78.34	*	-	
	HEMBA1003330	6.01	12.33	12.91	7.65	8.96	7.41	6.19	2.7	6.56			
	HEMBA1003348	3.04	4.46	6.33	4.48	9.58	12.27	5.11	0.11	2.09			
	HEMBA1003369	4.61	6.54	9.12	7.83	6.84	2.91	5.15	1.7	5.2			
45	HEMBA1003370	3.59	5.98	8.58	5.67	3.43	4.27	2.86	5.19	3.68			
	HEMBA1003373	11.8	17.95	18.19	5.05	9.3	7.7	3.82	6.22	4.16	*	**	-
	HEMBA1003376	3.97	3.38	4.86	3.82	1.87	2.14	1.54	0.79	1.25	**	-	
	HEMBA1003380	1.93	2.67	5.1	1.31	0.26	1.37	0.83	1.72	0.69			
	HEMBA1003384	26.49	38.34	57.87	29.22	29.1	27.24	18.27	17.42	21.24			
	HEMBA1003387	16.13	19.1	24.1	17.73	14.37	13.92	9.34	7.52	11.3	*	-	
50	HEMBA1003392	2.68	3.97	8.74	2.76	4.49	2.32	2.44	0.39	3.04			
	HEMBA1003395	2.69	3.42	9.32	1.54	4.93	3.86	3.02	1.02	1.64			
	HEMBA1003399	20.63	29.8	52.54	29.45	22.45	24.07	12.86	13.2	21.21			
	HEMBA1003400	3.56	5.58	6.33	3.88	4.46	4.23	1.87	2.25	1.87	*	-	
	HEMBA1003402	38.86	90.8	85.59	22.6	36.87	47.3	19.41	16.83	28.09	*	-	
	HEMBA1003403	4.18	3.19	7.45	2.51	2.47	2.16	1.4	2.51	3.27			
55	HEMBA1003408	25.83	33.63	38.04	42.18	33.38	41.52	23.7	19.86	28.4			
	HEMBA1003412	5.21	6.25	13.71	2.82	5	4.22	2.53	1.53	2.45			
	HEMBA1003417	21.02	35.52	34.24	13.8	14.17	10.79	5.28	6.88	9.43	*	**	-

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	HEMBA1002921	10.08	17.69	26.78	4.57	11.52	17.13	10.57	6.89	8.36			
	HEMBA1002924	16.52	23.06	41.44	17.55	21.51	18.51	16.69	6.29	12.01			
5	HEMBA1002934	7.43	7.93	15.24	8.33	19.71	10.41	9.2	4.34	7.83			
	HEMBA1002935	44.38	58.85	52.49	15.89	32.69	26.1	15.32	13.88	12.84	*	**	-
	HEMBA1002937	2.13	5.21	6.49	6.16	3.25	2.49	3.22	1.79	2.68			
	HEMBA1002939	6.4	9.23	9.9	6.39	7.51	3.18	2.69	1.8	1.53	**		-
	HEMBA1002944	6.74	7.31	12.98	7.18	5.53	7.34	7.96	2.7	9.59			
	HEMBA1002951	2.29	3.34	4.29	3.26	3.2	3.99	3	1.79	2.41			
10	HEMBA1002954	17.42	19.44	25.7	12.87	16.68	13.48	13.76	7.82	14.9			
	HEMBA1002962	8.79	11.09	12.46	11.77	11.02	6.43	11.04	6.23	10.79			
	HEMBA1002968	7.03	12.43	15.52	4.99	4.92	6.17	4.89	4.54	3.33	*		-
	HEMBA1002970	3.97	3.94	8.85	9.45	5.39	4.2	3.04	1.94	3.66			
	HEMBA1002971	46.35	65.62	65.66	55.83	32.32	29.43	18.21	22.18	16.03	**		-
	HEMBA1002973	19.1	22.88	32.02	18.54	14.79	14.79	14.19	10.32	16.42			
15	HEMBA1002978	9.61	14.49	12.56	9.49	10.35	10.49	8.97	6.63	9.32			
	HEMBA1002981	7.62	7.65	10.48	7.25	7.54	7.91	5.93	4.75	6.33			
	HEMBA1002985	117.94	141.29	115.79	100.57	91.73	67.47	36.07	32.99	67.86	*	**	-
	HEMBA1002986	1.09	4.19	2.41	1.9	4.92	2.49	2.41	1.95	2.68			
	HEMBA1002988	6.76	12.45	10.76	6.53	13.58	5.88	8.35	4.66	5.88			
20	HEMBA1002992	6.35	10.06	11.88	10.23	8.86	4.79	4.31	25.89	6.31			
	HEMBA1002995	4.3	8.89	7.73	5.51	4.55	4.85	3.59	3.01	4.66			
	HEMBA1002997	107.63	114.87	183.84	177.35	156.59	181.54	82.75	91.36	112.69			
	HEMBA1002999	53.34	68.01	72.24	37.39	47.67	41.25	62.98	43.59	63.81	*		-
	HEMBA1003004	3.64	4.33	7.42	2.58	3.67	3.65	3	1.57	3.06			
	HEMBA1003006	4.28	5.05	7.7	1.79	6.77	4.05	4.37	2.61	3.52			
25	HEMBA1003008	27.15	36.55	38.8	26.92	32.63	23.56	33.58	25.15	30.26			
	HEMBA1003021	5.81	12.5	13.45	11.13	6.12	5.84	7.32	6.94	6.02			
	HEMBA1003027	18.79	26.5	31.66	21.25	18.71	19.18	16.11	14.87	19.02			
	HEMBA1003029	19.01	29.05	31.26	23.38	17.83	18.43	16.34	15.3	19.09			
	HEMBA1003031	23.27	32.76	35.52	18.12	20.06	19.05	22.57	22.89	27.58	*		-
	HEMBA1003032	10.51	18.48	23.57	10.46	10.87	8.93	13.71	10.94	13.4			
30	HEMBA1003033	3.28	3.5	9.21	2.58	2.99	3.45	2.32	2.32	3.69			
	HEMBA1003034	32.22	43.17	44.46	25.3	23.69	20.88	30.34	23.98	27.83	*	*	-
	HEMBA1003036	6.89	6.34	9.49	7.01	5.99	4.57	6.73	1.95	2.1			
	HEMBA1003037	3.15	4.76	8.43	6.95	3.5	6.33	1.17	2.19	4.48			
	HEMBA1003041	5.81	4.77	6.88	6.46	7.11	3.98	3.65	3.49	6.68			
	HEMBA1003045	3.59	5.57	7.67	2.55	3.97	2.74	1.99	1	3.62			
35	HEMBA1003047	2.75	3.6	7.28	2.24	3.39	2.22	1.7	2.72	4.34			
	HEMBA1003048	7.72	9.91	17.38	5.16	6.98	5.24	7.18	5.24	6.81			
	HEMBA1003064	7.42	7.15	11.99	6.45	5.86	2.65	4.01	3.53	5.26	*		-
	HEMBA1003067	3.5	4.14	10.69	1.88	3.75	2.7	1.84	2.09	2.56			
	HEMBA1003071	4.3	6.5	8.99	2.07	3.4	1.57	3.49	0.09	2.65	*		-
	HEMBA1003072	22.65	30.56	72.67	40.85	32.84	36.51	17.64	22.4	33.19			
40	HEMBA1003076	10.41	7.13	12.54	12.87	5.59	14.57	5.56	3.38	7.99			
	HEMBA1003077	57.77	42.56	38.04	15.22	22.46	42.9	9.06	21.41	25.7	*		-
	HEMBA1003078	5.9	3.99	8.44	5.05	5.47	3.08	3.19	1.13	4			
	HEMBA1003079	2.67	1.28	11.95	1.19	1.44	0	1.26	1.15	2.5			
	HEMBA1003083	17.29	19.57	26.48	12.28	16.41	11.4	17.96	11.77	14.23			
	HEMBA1003086	66.37	81.42	75.05	81.77	61.06	81.5	45.88	45.17	41.59	**		-
45	HEMBA1003090	27.88	31.13	33.36	17.15	38.66	20.27	13.44	24.06	7.13	*		-
	HEMBA1003094	7.62	15.14	35.58	19.46	14.45	10.58	6.94	6.32	13.5			
	HEMBA1003096	5.15	5.97	7.99	4.67	5.25	3.62	2.44	1.23	5.06			
	HEMBA1003098	51.24	56.94	73.88	44.19	37.83	35.41	49.21	42.96	49.32	*		-
	HEMBA1003101	55.72	62.33	39.5	28.21	27.13	36.8	19.35	33.07	15.18	*	*	-
	HEMBA1003109	21.19	29.16	38.82	18.38	20.48	14.86	17.89	17.36	21.07			
50	HEMBA1003114	16.2	22.18	27.85	11.32	13.05	11.32	15.03	12.24	12.98	*		-
	HEMBA1003117	4.55	5.29	16.23	3.16	6.98	2.06	3.55	1.47	2.05			
	HEMBA1003120	26.44	23.65	42.74	11.15	27.45	17.66	13.62	6.31	12.11	*		-
	HEMBA1003129	13.18	14.01	38.23	23.62	19.83	17.41	9	7.3	18.68			
	HEMBA1003133	18.25	22.53	34.22	24.55	19.61	15.34	21.94	16.17	21.16			
	HEMBA1003136	16.39	19.51	25.59	18.11	18.16	15.75	14.64	13.22	14.96			
55	HEMBA1003142	14.56	16.27	15.48	9.65	8.16	8.79	8.14	5.26	6.82	**	**	-
	HEMBA1003148	16.81	26.84	29.42	16.45	15.92	11.14	21.43	11.51	11.59			
	HEMBA1003151	8.82	10.43	18.65	5.42	8.75	3.56	10.89	7.34	8.14			

Table 364

	HEMBA1002590	5.95	11.3	17.91	3.86	9.68	7.63	4.92	3.2	3.29			
	HEMBA1002592	24.56	35.63	62.38	31.85	31.39	28.3	31.44	17.58	25.5			
5	HEMBA1002595	18.13	24.94	28.1	20.67	22.43	16.85	23.31	9.38	10.88			
	HEMBA1002609	10.62	12.63	20.23	14.42	14.63	14.25	5.32	5.93	14.35			
	HEMBA1002617	4.27	5.73	7.4	1.57	2.4	3.96	1.42	3.15	4.02			
	HEMBA1002619	2.85	2.62	7.5	3.1	2.95	3.67	1.94	0.73	2.61			
	HEMBA1002621	8.59	8.17	23.61	10.11	9.98	16.3	7.86	5.98	9.09			
	HEMBA1002624	27.35	35.73	38.2	18.75	33.76	30.42	15.63	13.42	13.94	**	-	
10	HEMBA1002628	9.91	18.29	22.67	1.48	9.73	6.58	2.84	2.68	1.86	*	-	
	HEMBA1002629	17.6	28.89	36.07	15.14	20.64	15.47	19.33	12.69	24.63	*	-	
	HEMBA1002632	12.9	25.94	21.19	10.8	16.54	18.64	7.96	5.15	6.22	*	-	
	HEMBA1002645	10.16	17.89	28.87	12.29	14.08	14.33	9.35	8.1	11.96			
	HEMBA1002651	13.26	18.44	26.72	15.43	10.74	15.26	12.68	8.84	17.98			
15	HEMBA1002652	38.73	48.42	65.53	46.16	37.25	33.78	28.26	20.1	36.36			
	HEMBA1002659	9.03	11.88	19.16	8.75	10.05	12.88	11.02	6.8	9.75			
	HEMBA1002661	5.06	3.99	9.25	2.67	2.42	2.71	2.5	0.54	3.81			
	HEMBA1002666	4.98	2.76	10.31	1.02	2.15	2.47	1.95	1.39	2.97			
	HEMBA1002667	66.26	92.17	95.25	40.2	49.75	38.64	17.08	15.46	18.87	*	**	-
	HEMBA1002673	10.24	11.61	24.1	10.91	20.39	11.76	14.29	6.19	10.4			
20	HEMBA1002678	13.62	19.9	33.62	20.45	17.4	13.57	11.46	10.78	12.98	*	**	-
	HEMBA1002679	13.69	14.71	16.04	4.7	9.65	9.85	3.39	2.88	4.57	*	**	-
	HEMBA1002688	5.6	5.12	9.48	4.59	5.44	5.04	4.69	4.1	9.84			
	HEMBA1002696	59.06	73.44	43.41	32.21	38.96	30.65	22.83	22.73	43.71			
	HEMBA1002703	7.17	12.58	16.11	7.07	6.09	6.51	6.44	2.28	5.09			
	HEMBA1002706	6.13	6.22	16.32	2.59	4.89	3.24	4.44	2.11	3.85			
25	HEMBA1002712	11.08	15.22	21.38	7.53	9.92	6.72	6.23	2.15	5.09	*	-	
	HEMBA1002715	7.25	9.24	15.63	3.34	13.16	4.85	8.59	3.19	5.46			
	HEMBA1002716	10.24	5.79	8.57	7.72	6.63	11.75	6.68	8.82	8.25	*	-	
	HEMBA1002718	5.54	9.62	8.18	5.1	4.05	3.87	4.38	2.74	1.77	*	-	
	HEMBA1002728	12.44	14.25	13.51	6.78	9.73	5.54	5.16	3.78	4.32	*	**	-
	HEMBA1002730	18.89	21.85	36.97	19.48	14.69	13.56	21.24	13.64	21.33			
30	HEMBA1002734	18.2	30.15	26.54	25.61	18.78	20.96	17.72	11.44	17.14			
	HEMBA1002742	3.87	2.34	11.09	2.03	2.86	2.81	1.76	0.66	1.41			
	HEMBA1002746	22.55	32.65	52.23	26.89	25.86	22.2	22.76	13.04	18.94			
	HEMBA1002748	9.65	15.62	22.06	9.53	12.94	6.47	8.64	2.48	4.44			
	HEMBA1002750	4.66	4.65	9.25	7.27	4.67	4.79	3.27	2.62	3.14			
	HEMBA1002755	6.97	7.74	9.25	6.37	7.77	4.16	3.95	4.22	4.36	**	-	
35	HEMBA1002759	16.7	17.27	27.62	16.79	11.68	13.5	9.91	9.05	10.68	*	-	
	HEMBA1002763	47.58	57.94	56.78	54.23	36.74	38.37	36.64	19.12	21.65	*	-	
	HEMBA1002767	8.72	14.41	17.03	5.04	11.17	4.96	8.31	3.05	8.44			
	HEMBA1002768	5.52	5.94	12.79	2.88	4.3	2.49	2.51	1.17	1.65			
	HEMBA1002769	17.19	27.21	15.86	8.55	10.83	3.03	6.22	1.99	5.25	*	*	-
	HEMBA1002770	2.13	4.12	10.85	3.83	4.73	3.88	2.44	1.5	3.81			
40	HEMBA1002777	6.28	5.97	10	5.84	6.03	4.58	3.3	3.36	7.09			
	HEMBA1002779	4.32	3.05	9.05	3.43	3.09	3.54	2.97	2.09	4.26			
	HEMBA1002780	18.15	7.73	29.43	3.93	22.04	8.61	11.74	14.31	4.26			
	HEMBA1002790	44.22	50.48	73.09	31.77	35.83	33.38	43.74	34.78	30.04			
	HEMBA1002794	7.76	9.01	14.22	4.88	6.33	4.1	3.98	2.38	3.09	*	-	
	HEMBA1002798	58.19	53	86.62	94.91	76.86	71.68	49.35	25.85	43.83			
45	HEMBA1002801	3.51	8.44	16.4	3.59	4.66	6.72	3.9	1.17	4.1			
	HEMBA1002810	3.05	9.57	15.59	8.92	5.81	6.43	5.04	3.55	5.63			
	HEMBA1002816	33.54	55.39	45.57	5.42	16.08	37.06	10.38	10.74	17.79	**	-	
	HEMBA1002818	13.01	14.09	20.64	7.78	6.78	5.49	5.04	2.96	3.82	*	**	-
	HEMBA1002820	27.01	29.21	56.88	35.9	30.38	30.6	30.02	21.79	27.81			
	HEMBA1002826	19.2	24.63	59.75	28.13	24.2	21.9	24.8	13.89	24.13			
50	HEMBA1002833	63.25	54.77	137.72	30.62	26.06	40.67	37.5	33.73	24.96			
	HEMBA1002850	32.24	50.44	106.97	13.72	14.84	20.17	21.65	20.43	7.04			
	HEMBA1002862	2.91	4.02	13.2	2.77	4.61	2.19	2.67	1.11	1.52			
	HEMBA1002863	5.54	4.99	10.17	2.83	5.36	5.71	5.41	0.6	2.81			
	HEMBA1002867	12.64	16.67	40.23	22.98	17.46	25	12.79	7.67	12.96			
	HEMBA1002876	33.33	40.68	56.24	16.95	25.21	41.71	19.36	18.11	7.35	*	-	
55	HEMBA1002886	38.07	46.59	100.23	134.31	130.1	87.32	31.47	23.92	40.5			
	HEMBA1002896	373.7	87.77	442.56	153.27	289.24	233.48	153.73	231.71	129.4			
	HEMBA1002913	5.91	13.06	17.5	7.91	16.67	18.64	16.24	3.53	12.2			

Table 363

	HENBA1002356	36	42.3	70.07	29.6	29.36	25.6	38.93	22.92	26.48			
	HENBA1002357	5.52	13.1	13.45	6.21	11.96	5.72	6.55	2.43	6.78			
5	HENBA1002360	7.18	14.99	31.64	16.74	12.78	10.39	10.06	6.73	9.23			
	HENBA1002363	7.21	6.56	15.23	7.32	10.07	12.95	6.94	4.08	5.41			
	HENBA1002365	5.13	6.37	6.27	3.06	3.21	2.92	4.54	1.5	1.04	**	*	-
	HENBA1002370	2.68	3.2	4.12	1.48	1.53	1.16	1.82	1.08	0.59	*	*	-
	HENBA1002374	7.98	8.73	12.15	4.74	5.3	4.18	4.71	3.04	5.14	*	*	-
10	HENBA1002376	3.93	7.64	11.22	5.3	5.08	2.77	3.34	1	2.5			
	HENBA1002377	9.59	9.89	16.96	5.63	7.09	6.15	6.12	1.21	4.87	*		-
	HENBA1002380	10.33	16.4	28.08	14.11	13.73	11.4	15.31	7.9	14.99			
	HENBA1002381	4.05	9.29	15.58	7.37	6.4	6.25	6.86	1.71	6.19			
	HENBA1002384	8.47	11.56	16.49	7.99	8.37	10.32	10.3	5.34	12.08			
	HENBA1002389	11.42	20.29	29.05	5.75	14.68	8.45	8.49	15.83	6.08			
15	HENBA1002396	12.69	16.26	27.27	7.41	9.5	6.56	5.14	3.17	3.81	*		-
	HENBA1002402	11.74	16.83	21.99	13.15	10.97	7.11	8.38	8.96	10.5			
	HENBA1002417	16.84	19.38	28.25	14.12	13.6	10.71	18.74	11.52	12.94			
	HENBA1002419	5.76	8.38	11.36	2.24	4.5	1.91	3.6	1.74	2.93	*	*	-
	HENBA1002420	19.24	23.07	51.09	26.34	25.11	18.31	20.11	12.19	18.92			
	HENBA1002421	18.39	32.76	48.68	28.36	38.24	25	28.96	7.94	14.32			
20	HENBA1002423	22.61	32.68	44.06	7.09	7.86	5.55	4.56	5.5	6.54	*	*	-
	HENBA1002424	10.97	10.88	26.72	15.1	10.87	9.61	12.1	6.14	11.48			
	HENBA1002426	7.39	16.83	33.68	10.19	19.75	11.23	14.09	9.36	9.51			
	HENBA1002430	3.4	2.99	7.25	1.05	5.01	3.49	1.89	3.82	1.55			
	HENBA1002439	16.88	14.68	38.41	5.15	12.09	6.57	10.43	3.04	5.71			
25	HENBA1002441	6.85	11.02	20	8.08	6.54	5.94	8.36	4.52	6.62			
	HENBA1002454	12.57	18.57	41.83	19.84	16.12	15.91	15.68	8.28	16.15			
	HENBA1002458	18.76	29.12	42.05	23.99	29.98	19.78	21.68	9.8	15.93			
	HENBA1002460	17.05	22.65	60.16	30.55	31.36	38.17	25.74	12.29	25.18			
	HENBA1002462	4.29	4.96	6.67	4.04	2.82	5.54	6.09	4.59	1.34			
	HENBA1002465	18.46	17.14	33.42	10.69	9.15	14.44	4.41	4.25	2.08	*		-
	HENBA1002469	5.68	18.77	19	7.83	8.55	8.09	9.48	5.61	6.29			
30	HENBA1002475	4.8	4.95	7.67	3.53	3.61	2.82	2.16	0.66	3.66	*		-
	HENBA1002477	14.5	19.14	47.19	10.6	19.29	17.91	11.51	6.53	9.54			
	HENBA1002480	3.04	4.91	11.31	4.99	4.91	2.26	3.45	1.28	2.5			
	HENBA1002481	3.73	5.8	10.36	6.68	9.38	9.35	6.55	0.82	5.73			
	HENBA1002486	14.96	20.73	36.85	17.9	13.84	17.9	9.8	9.17	15.34			
35	HENBA1002490	32.89	43.68	75.36	48.87	29.68	38.22	19.18	16.64	20.65			
	HENBA1002495	43.66	58.23	82.55	111.1	32.51	41.26	76.83	29.1	14.88			
	HENBA1002498	3.78	6.46	9.54	6.71	5.27	12.7	10.64	3.53	6.2			
	HENBA1002501	7.77	5.26	15.06	10.61	8.44	10.8	6.66	2.76	7.39			
	HENBA1002503	1.3	3.76	4.26	11.14	4.16	3.06	2.94	0.74	2.46			
	HENBA1002504	11.32	6.35	15.09	5.12	7.76	6.01	3.02	1.3	4.4	*		-
	HENBA1002508	5.06	9.32	4.01	3.71	7.02	5.17	4.27	2.18	3.39			
40	HENBA1002513	9.71	15.43	17.71	11.5	8.94	8.42	3.65	5.7	5.29	*		-
	HENBA1002515	22.59	29.58	33.08	19.66	17.58	7.29	3.31	4.58	5.25	**		-
	HENBA1002524	2.52	5.5	6.81	5.24	5.07	5.47	3.51	2.04	6.31			
	HENBA1002538	4.86	8.27	13.28	5.54	6.86	8.78	7.76	3.44	3.36			
	HENBA1002542	14.29	22.48	34.17	23.25	22.99	20.23	26.21	13.64	24.16			
45	HENBA1002544	80.29	71.81	142.33	191.63	103.72	120.69	75.35	42.62	73.64			
	HENBA1002546	16.49	22.78	27.12	14.78	15.43	17.06	9.62	3.52	9.44	*		-
	HENBA1002547	3.95	4.98	8.23	6.73	9.06	6.15	8.5	2.36	7.03			
	HENBA1002550	49.15	78.71	90.52	40.75	20.92	26.29	11.49	22.91	24.84	*	*	-
	HENBA1002551	21.39	26.41	43.07	27.63	15.98	20.86	15.81	13.41	18.29			
	HENBA1002552	2.51	5.57	9.92	2.51	4.26	4.71	1.72	3.19	4.1			
	HENBA1002555	22.2	20.07	42.05	23.19	22.23	22.76	15.94	12.91	13.88			
50	HENBA1002558	12.09	15.97	19.47	12.44	11.51	11.85	16.99	12.78	19.34			
	HENBA1002561	5.12	5.08	10.95	9.59	4.94	7.59	3.52	2.9	5.93			
	HENBA1002562	11.02	16.11	24.74	19.26	26.78	27.85	11.51	7.07	19.83			
	HENBA1002568	196.94	317.85	248.99	300.72	297.51	250.18	199.37	60.64	136.04			
	HENBA1002569	23.5	41.31	56.08	12.56	22.81	12.46	7.29	13.02	18.03			
	HENBA1002570	30.8	40.14	52.73	26.65	25.92	32.98	19.03	21.81	22.76	*		-
55	HENBA1002574	18.81	26.29	36.64	14.11	18.09	18.3	12.06	11.37	15.9			
	HENBA1002582	7.94	6.71	16.38	5.31	5.66	5.07	2.73	4.44	3.12			
	HENBA1002587	2.94	4.41	12.22	7.72	5.53	9.69	4.6	2.49	5.42			

Table 362

	HEMBA1002133	29.03	35.39	76.13	107.6	70.61	84.67	29.03	14.5	31.23		
	HEMBA1002139	11.59	23.7	52.84	28.04	21.84	20.31	32.1	12.89	25.76		
5	HEMBA1002141	3.72	5.77	10.55	5.21	5.75	3.18	3.83	1.34	5.31		
	HEMBA1002144	2.85	4.6	6.42	4.94	2.2	5.21	1.49	2.58	2.66		
	HEMBA1002147	10.27	15.44	23.81	11.25	8.38	13.35	8.76	6.54	9.55		
	HEMBA1002150	1.33	1.23	3.53	2.46	2.28	2.8	1.93	0.73	2.03		
	HEMBA1002151	6.03	4.63	11.41	5.99	5.56	6.68	5.34	2.78	4.91		
	HEMBA1002153	0.77	2.83	4.03	3.46	4.19	3.78	2.71	1.39	3.61		
10	HEMBA1002156	2.06	3.91	4.83	5.6	4.46	2.69	2.87	0.76	2.49		
	HEMBA1002160	56.44	32.19	84.15	108.2	70.78	79.86	43.2	16.4	44.96		
	HEMBA1002161	44.7	21.27	41.93	61.65	56.83	51.89	26.74	12.83	24.21		
	HEMBA1002162	89.86	103.46	226.68	313.25	143.99	221.76	91.22	73.96	98.59		
	HEMBA1002163	40.99	33.35	79.59	79.2	53.49	69.59	30.86	22.17	36.21		
	HEMBA1002164	5.75	8.38	13.54	8.04	3.83	5.52	3.45	2.03	3.62		
15	HEMBA1002166	15.02	15.38	25.99	20.34	17.99	15.76	15	11.24	18.28		
	HEMBA1002167	11.24	16.13	23.3	10.71	13.78	13.32	12.46	7.17	15.43		
	HEMBA1002173	2.74	4.74	7.08	7.8	4.98	8.26	4.32	0.11	6.96		
	HEMBA1002177	3.19	7.92	6.73	9.46	13.68	11.32	3.6	0.89	5.89	*	+
	HEMBA1002178	3.07	3.14	5.24	4.35	5.65	4.54	2.7	1.64	3.18		
	HEMBA1002179	4.9	4.72	9.91	6.66	4.93	4.53	2.62	2.86	3.55		
20	HEMBA1002185	24.67	32.27	51.89	14.08	21.43	18.77	8.38	11.95	12.92	*	-
	HEMBA1002188	5.88	6.14	7.11	2.97	4.39	3.33	2.27	4.54	2.68	**	-
	HEMBA1002189	69.59	81.69	157.25	233.85	141.07	152.77	75.89	48.73	66.24		
	HEMBA1002191	4.08	3	7.66	6.05	7	4.24	3.01	1.76	4.2		
	HEMBA1002192	5.95	7	11.91	7.01	6.79	6.52	4.64	2.39	5.33		
25	HEMBA1002195	4.46	6.6	10.03	6.4	6.23	6.42	6.4	2.77	4.74		
	HEMBA1002196	11.48	14.42	16.69	14.62	12.56	12.05	7.65	1.78	5.06	*	-
	HEMBA1002199	17.19	29.45	44.34	8.12	15.7	10.6	4.22	6.56	8.91	*	-
	HEMBA1002204	12.36	17.29	25.36	18.54	15.65	13.06	6.95	7.55	12.67		
	HEMBA1002208	4.42	3.1	8.67	2.76	2.3	3.61	1.67	2.17	2.81		
	HEMBA1002212	8.54	9.41	18.06	5.8	6.49	9.16	8.07	6.31	7.72		
	HEMBA1002215	3.77	4.75	11.1	3.82	3.64	4.93	3.19	2.31	3.82		
30	HEMBA1002217	3.21	2.97	8.05	1.88	2.5	3.26	2.77	1.7	3.42		
	HEMBA1002220	40.07	58.29	38.41	33.44	37.1	26.5	27.12	21.87	26.71	*	-
	HEMBA1002226	3.86	6.37	9.01	5.94	8.12	6.13	6.11	1.94	2.28		
	HEMBA1002227	29.2	48.92	80.06	21.26	24.93	28.49	9.06	21.04	35.76		
	HEMBA1002229	33.53	65.16	51.1	12.99	38.46	30.19	10.92	21.02	13.32	*	-
35	HEMBA1002237	5.24	7.25	13.43	6.25	5.49	6.14	5.27	3.9	7.13		
	HEMBA1002239	4.63	2.53	7.61	2.56	2.45	2.52	2.39	1.83	5.54		
	HEMBA1002241	106.58	86.21	101.41	191.32	99.49	119.15	95.67	52.74	66.24		
	HEMBA1002253	12.22	17.16	20.51	2.71	7.33	6.42	3.87	2.39	2.33	*	**
	HEMBA1002257	7.54	10.24	15.29	4.58	6.87	7.36	4.9	4.18	3.85	*	-
	HEMBA1002259	13.5	22.26	29.8	14.68	19.62	16.23	17.99	11.27	18.91		
40	HEMBA1002262	15.09	23.24	29.3	18.72	15.42	11.55	9.03	9.95	14.69		
	HEMBA1002265	14.18	19.86	21.44	12.15	13.48	12.48	8.23	10.18	13.41	*	-
	HEMBA1002267	27.68	15.47	50.07	10.16	19.76	26.78	8.51	9.81	8.8		
	HEMBA1002270	6.74	5	13	3.57	4.76	3.79	3.06	1.88	3.1		
	HEMBA1002286	3.26	3.45	8.41	1.32	1.98	2.17	2.49	0.44	3.13		
	HEMBA1002290	13.88	14.35	29.79	17.54	14.92	16.25	8.98	7.54	11.64		
45	HEMBA1002302	2.59	3.21	9.4	1.67	2.25	2.2	2.59	2.03	2.91		
	HEMBA1002304	28.75	42.22	44.71	47.37	56.93	39.52	32.89	19.02	22.8		
	HEMBA1002307	11.61	12.1	29.02	16.8	17.62	14.8	7.63	7.75	11.14		
	HEMBA1002316	59.64	53.59	76.81	100.13	70.41	86.41	40.44	34.11	39.43	*	-
	HEMBA1002319	5.85	5.03	12.11	6.62	5.15	5.86	4.45	4.18	4.84		
	HEMBA1002320	5.15	6.41	11.19	4.43	3.26	5.68	2.72	1.07	1.82	*	-
50	HEMBA1002321	26.14	32.72	45.81	15.36	15.91	12.15	25.74	14.29	26.13	*	-
	HEMBA1002328	3.49	3.33	9.66	1.53	2.94	1.89	1.67	1.15	3.51		
	HEMBA1002333	14.03	17.53	33.87	10.39	17.53	10.54	14.85	12.2	18.17		
	HEMBA1002337	74.1	90.11	104.64	39.01	88.83	67.57	65.51	46.31	39.48	*	-
	HEMBA1002339	7.99	5.95	12.61	11.75	8.3	7.98	3.81	2.17	4.39		
	HEMBA1002341	10.07	14.65	15.74	8.32	6.89	8.79	4.57	2.09	3.76	*	**
55	HEMBA1002348	59.9	42.49	67.67	22.81	38.26	47.94	42.75	47.55	14.2		
	HEMBA1002349	3.79	4.77	8.57	3.9	2.41	2.17	3.1	1.47	1.4		
	HEMBA1002353	16.84	19.17	38.76	16.44	19.61	18.36	31.34	14.04	19.17		

Table 361

	HENBA1001879	23.8	24.57	36.17	27.25	30.71	21.12	18.92	12.59	21.51			
	HENBA1001884	22.79	25.41	39.57	18.36	21.7	20.15	24.74	18.16	27.02			
5	HENBA1001886	2.49	3.21	10.85	4.21	4.2	3.06	3.99	2.94	2.81			
	HENBA1001888	5.93	5.79	13.13	8.79	5.21	8.71	5.32	6.52	7.9			
	HENBA1001890	12.55	17.2	16.26	7.93	7.84	9.91	5.34	7.04	9.79	*	*	-
	HENBA1001896	8.62	13.4	18.92	9.94	9.97	13.19	7.37	7.12	11.57			
	HENBA1001899	9.63	9.32	14.65	9.07	9.12	9.82	4.74	3.78	7.78			
	HENBA1001904	17.69	9.43	25.81	8.49	15.82	16.23	7.42	3.21	4			
10	HENBA1001910	13.84	17.03	18.88	10.76	16.22	11.5	15.72	11.44	11.3			
	HENBA1001911	84.18	136.96	104.54	60.48	92.99	36.82	38.97	44.93	16.51	*		-
	HENBA1001912	8.8	18.15	15.1	11.38	13.1	11.76	6.02	5.45	6.48	*		-
	HENBA1001913	63.5	67.85	178.77	91.97	76.48	71.89	38.1	40.9	79.58			
	HENBA1001915	9.95	12.39	24.42	17.53	14.5	13	8.17	4.57	10.88			
	HENBA1001918	5.57	6.46	12.5	8.11	6.81	8.09	4.3	2.18	7.19			
15	HENBA1001921	8.68	9.91	12.62	8.57	8.25	7.86	5.28	3.66	4.78	**		-
	HENBA1001931	9.24	10.47	21.38	7.16	7.58	6.86	8.41	6.87	7.96			
	HENBA1001939	3.59	7.32	15.17	1.25	16.55	5.08	4.84	5.03	14.82			
	HENBA1001940	11.2	9.93	21.45	9.87	12.2	10.55	7.02	5.09	7.03			
	HENBA1001942	22.25	36.8	27.34	8.22	35.32	31.27	17.26	9.8	12.44	*		-
20	HENBA1001944	14.23	17.39	30.07	21.25	14.66	13.47	4.56	8.32	14.97			
	HENBA1001945	8.53	9.04	18.07	9.58	9.66	6.81	6.7	4.7	10.21			
	HENBA1001950	35.24	36.06	25.81	22.39	20.56	21.76	31.55	22.6	14.77	*		-
	HENBA1001951	39.37	38.17	52.67	44.58	33.39	34.05	41.69	23.81	45.03			
	HENBA1001958	3.68	3.88	9.13	1.87	2.36	1.68	2.36	1.43	3.59			
	HENBA1001960	11.92	13.79	24.84	7.91	13.38	11.06	7.97	3.56	9.09			
25	HENBA1001962	10.82	11.31	28.75	9.3	13.76	9.37	10.95	7.24	13.69			
	HENBA1001964	11.99	14.1	28.18	9.47	13.95	12.22	12.21	4.77	10.5			
	HENBA1001967	33.16	31.11	63.39	22.3	13.19	20.77	8.29	10.21	24.27			
	HENBA1001979	6.77	6.74	6.61	3.38	4.61	6.79	4.27	2.37	4.75	*	*	-
	HENBA1001987	6.84	5.64	7.7	5.03	3.79	3.32	4.73	5.25	2.63	*		-
	HENBA1001991	4.14	2.8	6.6	1.78	2.7	1.18	2.19	1.73	1.57			
30	HENBA1002003	3.19	3.64	8.31	2.58	2.51	1.13	4.14	2.93	2.53			
	HENBA1002005	21.43	25.67	43.64	22.86	20.88	16.44	24.76	11.27	20.55			
	HENBA1002008	15.2	19.08	39.6	14.05	18.61	13.62	20.77	11.98	21.03			
	HENBA1002016	4.58	11.59	12.53	3.51	5.86	20.49	4.43	1.81	2.74			
	HENBA1002022	29.07	36.45	75.11	52.83	42.93	48.33	30.25	25.12	55.78			
	HENBA1002029	19.99	24.25	30.34	10.7	17.32	16.65	10.55	13.15	11.59	*	*	-
35	HENBA1002030	11.06	12.89	19.22	8.24	7.3	5.76	8	6.82	6.24	*	*	-
	HENBA1002035	3.07	17.04	7.87	2.25	5.11	2.16	4.64	3.69	6.06			
	HENBA1002037	95.17	141.19	140.82	59.83	84.82	73.74	61.45	42.2	28.14	*	*	-
	HENBA1002038	11.96	19.11	18.49	6.44	9.02	6.74	4.63	2.27	2.59	*	**	-
	HENBA1002039	6.42	7.41	13.49	2.37	5.91	2.48	3.27	1.37	3.16	*	*	-
	HENBA1002042	3.25	8.61	11.52	6.6	6.65	6.58	7.97	0.75	4.67			
40	HENBA1002043	13.24	16.58	22.47	5.13	6.87	6.08	5.01	5.72	8.43	*	*	-
	HENBA1002048	25.65	31.58	36.46	5.14	12.63	17.17	11.27	12.56	24.5	*	*	-
	HENBA1002049	431.22	564.03	858.02	234.02	525.06	482.28	304.41	585.23	233.29			
	HENBA1002053	66.35	61.58	114.43	104.42	98.85	104.79	56.5	40.65	69.44			
	HENBA1002055	19.16	20.77	33.11	22.38	16.8	17.16	15.07	11.5	15.77			
	HENBA1002056	5.26	6.55	10.96	2.52	2.93	4.72	5.3	2.16	5.03			
45	HENBA1002061	3.02	4.19	9.86	2.45	3.45	0.92	2.37	0	1.82			
	HENBA1002080	2.59	8.86	20.11	3.27	8.34	5.7	6.09	2.76	4.82			
	HENBA1002084	7.01	3.85	10.52	5.96	3.47	4.97	2.69	1.31	2.02			
	HENBA1002085	3.68	5.88	10.92	2.9	5.24	7.31	3.65	1.95	5			
	HENBA1002092	18.94	17.76	38.17	14.49	14.47	17.01	14.19	7.25	10.23			
	HENBA1002098	8.74	16.29	15.2	3.59	8.82	8.02	9.31	14.32	5.83			
50	HENBA1002100	13.69	10.18	19.02	11.56	6.49	4.52	8.34	7.69	7.11			
	HENBA1002101	12.62	14.09	31.93	14.79	12.76	11.05	16.17	8.68	13.96			
	HENBA1002102	4.18	7.54	13.28	4.1	4.71	4.56	3.77	3.14	3.09			
	HENBA1002105	5.07	7.15	15.22	5.77	3.71	9.62	4.21	1.98	2.79			
	HENBA1002107	7.22	7.4	21.27	9.53	10.18	12.18	2.67	2.93	5.58			
	HENBA1002113	5.14	7.17	14.68	10.73	7.63	8.56	6.5	2.35	7.99			
55	HENBA1002119	24.1	26.96	60	38.03	31.14	26.41	31.91	16.47	21.72			
	HENBA1002125	4.31	5.62	9.66	11.65	3.83	5.15	2.02	2.4	2.97			
	HENBA1002131	2612	1876.5	3959.7	1306.5	2879.6	1519.6	1255.8	1936.4	1651.4			

Table 360

	HEMBA1001636	3.01	3.66	7.63	2.63	2.16	1.96	2	2.43	2.09			
	HEMBA1001640	29.8	43.06	50.77	23.27	24.57	21.36	34.37	24.57	21.57	*	-	
5	HEMBA1001647	5.45	3.94	16.28	4.05	6.97	2.61	5.21	2.15	4.19			
	HEMBA1001651	40.26	49.2	70.61	64.68	55.88	51.2	30.59	17.33	15.12	*	-	
	HEMBA1001655	10.17	9.88	20.1	6.79	14.34	13.87	15	5.01	12.4			
	HEMBA1001658	5.48	21.34	12.68	3.21	27.79	21.94	17.32	4.91	4.48			
	HEMBA1001661	1.92	2.27	3.66	1.72	2.22	1.93	2.04	0.71	3.46			
	HEMBA1001665	5.27	5.54	9.55	3.17	2.96	2.46	2.94	1.87	2.02	*	*	-
10	HEMBA1001670	14.59	18.93	22.5	10.01	11.18	7.78	13.23	9.01	14.55	*	-	-
	HEMBA1001672	5.67	6.19	14.07	4.38	5.16	2.99	5.11	2.78	4.53			
	HEMBA1001673	67.5	83.13	100.8	79.52	76.6	62.66	79.54	46.76	80.49			
	HEMBA1001675	11.65	14.39	26.02	11.92	12.54	11.44	8.96	6.24	12.74			
	HEMBA1001676	5.94	6.55	16.33	9.03	9.2	9.7	5.97	3.49	7.17			
15	HEMBA1001678	408.84	460.84	545.99	212.4	385.11	245.37	243.82	358.42	324.52	*	*	-
	HEMBA1001680	17.81	21.9	34.13	4.77	16.15	9.85	5.24	7.39	4.29	*	-	-
	HEMBA1001681	4.07	7.09	7.63	4.19	6.03	3.95	5.56	3.95	9.27			
	HEMBA1001684	9.98	10.79	18.76	5.52	7.17	3.85	8.63	7.5	7.69			
	HEMBA1001695	6.3	10.78	12.26	4.59	5.86	5.01	6.15	3.07	8.56			
	HEMBA1001702	4.21	4.98	12.28	2.87	3.73	2.48	2.67	0.5	1.76			
	HEMBA1001709	2.14	4.26	13.24	2.55	2.4	2.49	3	0.8	2.49			
20	HEMBA1001711	4.46	2.9	7.77	4.65	5.15	6.37	2.22	1.03	3.22			
	HEMBA1001712	17.3	21.73	24.38	4.68	4.43	7.93	6.13	3.94	7.49	**	**	-
	HEMBA1001714	5.44	5.05	12.26	3.87	6.98	3.98	4.35	1.76	2.27			
	HEMBA1001717	12.02	13.24	20.14	10.04	12.58	8.86	8.97	5.33	8.75			
	HEMBA1001718	6.01	7.09	16.45	6.91	7.4	6.4	6.45	5.23	4.35			
25	HEMBA1001723	4.38	5.84	13.91	3.89	3.74	3.51	3.82	2.36	1.25			
	HEMBA1001731	2.72	5.88	12.37	2.09	2.33	1.98	2.59	1.24	1.73			
	HEMBA1001734	3.48	4.86	15.18	4.34	3.76	3.13	3.75	1.03	2.79			
	HEMBA1001736	7.38	10.86	24.38	12.76	10.32	13.2	4.98	2.9	5.27			
	HEMBA1001741	15.06	22.96	34.81	18.57	17.99	31.53	19.64	14.22	21.03			
	HEMBA1001744	3.63	4.77	7.03	5.75	5.45	4.88	3.08	0.38	1.67			
30	HEMBA1001745	6.61	5.99	11.87	4.17	5.23	3.45	2.53	2.9	1.77	*	-	
	HEMBA1001746	10	14.3	20.05	11.55	8.04	11.63	7.31	4.66	10.54			
	HEMBA1001761	4.09	4.47	15.81	2.95	5.47	4.97	4.55	2.51	3.77			
	HEMBA1001762	12.27	16.22	31.71	11.79	12.77	9.07	10.3	5.04	7.64			
	HEMBA1001781	15.21	11.25	22.91	6.13	9.86	7.74	12.65	6.92	10.68			
	HEMBA1001784	7.14	8.4	14.02	11.89	8.86	32.45	4.84	4.18	5.94			
35	HEMBA1001791	5.49	5.04	11.37	7.91	5.03	6.87	4.96	3.16	6.54			
	HEMBA1001794	20.29	19.03	24.02	20.07	12.65	17.58	18.71	4.14	12.87			
	HEMBA1001800	3.21	4.6	6.87	7.34	5.09	5.03	5.14	1.64	4.39			
	HEMBA1001803	13.17	17	29.4	25.23	15.99	17.41	14	9.49	14.18			
	HEMBA1001804	10.28	12.82	18.21	18.52	13.37	6.76	5.06	3.75	9.33			
	HEMBA1001808	1.39	1.46	0.57	2.77	3.18	1.86	0.98	1.2	0.86	*	+	
40	HEMBA1001809	10.03	14.07	29.16	27.06	14.98	19.9	10.22	6.76	12.88			
	HEMBA1001811	4.85	4.27	3.18	10.29	3.63	4.31	3.6	2.84	5.25			
	HEMBA1001815	12.35	11.27	24.93	14.76	10.57	16.04	11.75	6.51	11.06			
	HEMBA1001816	15.52	16.02	31.22	15.84	13.09	17.05	15.48	10.68	15.2			
	HEMBA1001819	6.04	14.01	18.63	8.04	9.05	17.69	7.85	3.75	9.93			
	HEMBA1001820	1.8	2.22	6.75	3.5	2.81	3.45	1.98	0.63	2.02			
45	HEMBA1001822	10	12.84	23.21	14.57	16.12	12.64	12.36	6.85	15.78			
	HEMBA1001824	5.2	8.63	7.51	6.46	8.51	5.78	4.67	0.98	4.7			
	HEMBA1001835	9.56	9.62	14.11	4.64	5.39	6.11	3.44	4.66	3.04	*	**	-
	HEMBA1001844	8.02	10.52	19.05	12.42	10.73	9.57	5.22	6.54	4.99			
	HEMBA1001847	5.48	6.03	7.01	5.38	4.93	4.88	3.96	3.19	3.69	**	-	
	HEMBA1001849	6.93	9.14	14.78	5.03	7.86	7.21	7.8	8.87	10.07			
50	HEMBA1001850	16.31	17.09	25.61	18.6	20.16	20.58	11.51	7.47	10.97	*	-	
	HEMBA1001861	2.36	3.42	4.47	4.07	3.37	3.99	2	1.11	3.64			
	HEMBA1001862	13.85	31.74	17.1	16.22	16.82	16.11	10.11	2.6	10.67			
	HEMBA1001864	2.79	3.87	7.65	5.32	5.64	4.4	3.29	2.25	6.43			
	HEMBA1001866	5.87	7.28	17.71	11.53	8.34	6.62	3.42	4.81	5.52			
	HEMBA1001869	12.96	19.53	27.66	23.68	16.72	23.14	5.91	10.06	12.62			
55	HEMBA1001871	3.56	6.35	8.61	4.54	2.2	3.49	3.21	1.17	2.75			
	HEMBA1001876	4.6	3.93	8.22	5.16	4.46	4.33	3.76	4.15	6.11			
	HEMBA1001878	8.8	14.72	27.92	7.18	13.52	10.62	6.43	4.37	4.82			

Table 359

	HEMBA1001364	5.59	5.71	8.25	5.34	8.32	6.23	4.19	2.87	3.41	*	-
	HEMBA1001375	10.82	8.04	9.96	6.56	6.52	8.55	3.93	8.03	4.02		
5	HEMBA1001377	30.09	30.45	47.44	34.01	31.38	34.79	26.34	19.08	22.43		
	HEMBA1001383	3.91	6.12	10.35	4.08	4.49	3.29	3.04	1.16	2.13		
	HEMBA1001387	4.6	4.17	9.82	4.35	3.61	5.27	5	2.3	4.29		
	HEMBA1001388	3.38	12.99	10.77	8.84	13.96	11.27	14.15	2.79	4.79		
	HEMBA1001390	16.62	20.96	41.58	20.9	20.32	29.35	22.75	17.7	27.56		
10	HEMBA1001391	2.64	5.63	6.55	6.35	4.1	4.57	3.4	1.38	3.14		
	HEMBA1001398	7.95	9	20.79	14.55	10.11	10.1	9.09	3.86	9.96		
	HEMBA1001405	12.88	15.37	29.38	17.27	15.08	16.39	13.42	7.68	15.76		
	HEMBA1001406	1.07	1.48	6.07	4.64	2.23	2.45	3.84	0.53	2.8		
	HEMBA1001407	26.07	42.46	69.39	34.3	28.94	28.07	26.15	15.14	21.96		
	HEMBA1001411	12.19	10.95	23.76	12.34	12.91	16.6	13.01	4.8	11.91		
	HEMBA1001413	3.96	7.52	13.68	7.68	4.74	6.7	4.77	2.46	3.91		
15	HEMBA1001414	10.32	10.99	22.49	13.09	11.95	9.01	7.42	9.57	11.87		
	HEMBA1001415	108.07	112.24	218.31	263.58	139.52	200.62	94.4	74.64	97.27		
	HEMBA1001416	5.94	4.6	7.89	5.06	3.57	4.15	3.74	1.73	3.21	*	-
	HEMBA1001432	3.3	3.49	7.34	4.93	5.95	7.57	4.22	3.23	2.6		
	HEMBA1001433	5.11	5.55	7.48	5.92	7.1	6.22	4.04	2.29	5.52		
20	HEMBA1001435	448.61	559.24	661.15	777.73	679.52	650.29	640.12	377.73	869.06		
	HEMBA1001442	5.84	10.08	7.28	7.83	12.04	8.46	5.85	1.96	2.4		
	HEMBA1001448	3.77	5.34	6.49	8.2	4.64	4.7	3.45	2.22	3.65		
	HEMBA1001450	87.92	103.44	150.5	232.41	124.8	175.34	89.47	74.72	92.31		
	HEMBA1001454	32.58	33.13	59.47	19.79	12.7	25.66	18.88	24.9	33.22		
	HEMBA1001455	4.27	6.3	7.42	3.13	5.27	4.96	6.16	1.51	6.96		
25	HEMBA1001459	13.66	26.28	34.13	12.19	9.69	14.19	9.18	11.22	13.6		
	HEMBA1001461	30.98	26.88	43.29	63.45	32.06	36.73	26.62	18.12	23.29		
	HEMBA1001462	9.43	9.05	11.64	4.2	7.04	5.63	4.19	2.77	3.22	*	**
	HEMBA1001463	1.15	4.7	4.74	4.99	6.6	4.02	4.68	1.57	3.46		
	HEMBA1001469	23.94	42.87	64.76	64.17	33.38	52.34	14.36	18.57	20.01		
	HEMBA1001473	7.37	9.89	11.35	4.59	3.3	4.44	2.71	5.14	2.51	*	*
30	HEMBA1001477	143.72	139.98	283.29	91.65	145.02	139.05	125.85	114.31	63.92		
	HEMBA1001478	3.38	2.99	11.83	2.63	2.55	3.35	2.27	2.13	2.08		
	HEMBA1001480	12.29	13.63	23.83	4.88	9.2	12.63	3.97	2.72	4.12	*	-
	HEMBA1001483	20.97	27.37	23.32	22.22	18.94	28.05	14.33	10.01	19.51	*	-
	HEMBA1001490	14.01	20.37	28.13	15.26	17.45	15.48	22.44	13.02	12.6		
	HEMBA1001495	29.87	31.18	55.68	75.52	52.85	69.44	21.73	15.8	24.63		
35	HEMBA1001497	50.34	63.13	114.17	128.19	117.69	137.5	43.07	37.4	51.44		
	HEMBA1001510	6.77	7.43	13.73	8.69	8.1	7.82	5.34	3.9	10.87		
	HEMBA1001515	4.96	2.98	9.2	1.95	4	3.02	1.96	0.89	7.49		
	HEMBA1001517	8.97	7.98	15.08	6.1	6.94	5.43	7.39	5.25	8.47		
	HEMBA1001522	12.78	16.29	21.22	10.39	13.01	11	10.54	9.25	10.66		
	HEMBA1001526	8.05	10.86	20.31	6.19	10.35	7.24	6.93	4.75	7.32		
40	HEMBA1001533	10.67	13.52	23.12	12.09	16.84	9.55	9.42	6.16	11.58		
	HEMBA1001547	5.91	8.31	10.54	8.17	5.66	9.57	6.23	2.5	7.38		
	HEMBA1001552	2.97	4.72	5.75	6.36	5.56	2.49	2.68	1	5.09		
	HEMBA1001553	11.17	10.19	22.05	10.98	12.27	11.05	5.91	7.82	10.68		
	HEMBA1001557	7.49	2.45	11.09	3.83	4.84	5.27	3.14	6.97	8.01		
	HEMBA1001563	6.6	2.01	15.37	5.55	8.29	6.53	4.15	2	6.53		
45	HEMBA1001566	4.27	3.01	8.62	2.13	3.6	3.87	2.2	1.62	3.88		
	HEMBA1001569	9.15	15.35	27.87	11.41	11.78	14.08	8.02	5.27	11.03		
	HEMBA1001570	10.89	22.08	19.57	12.99	13.03	16.53	9.14	4.13	3.81	*	-
	HEMBA1001579	95.92	121.26	222.36	134.48	102.06	70.96	25.6	43.38	73.16		
	HEMBA1001581	4.11	4.25	7.71	3.25	2.45	2.33	0.91	1.64	3.55		
	HEMBA1001582	12.19	16.67	20.04	15.87	10.82	11.89	13.98	8.69	21.33		
50	HEMBA1001585	6.81	4.99	13.1	7.76	5	7.16	4.04	1.85	5.57		
	HEMBA1001589	7.95	8.96	19.02	7.86	6.17	6	7.81	5.42	8.45		
	HEMBA1001595	152.72	179.08	178.84	145.86	142.24	141.98	126.66	83.83	110.77	*	*
	HEMBA1001604	23.66	24.89	37.02	23.72	21.32	19.26	15.97	14.33	15.78	*	-
	HEMBA1001608	43.55	20.96	66.71	67.7	43.63	57.55	24.58	18.07	23.43		
	HEMBA1001615	5.42	4.63	6.88	6.61	4.72	4.09	3.11	2.02	10.18		
55	HEMBA1001620	30.69	48.66	53.49	25.04	17.81	22.67	7.35	9.47	10.42	*	**
	HEMBA1001621	8.82	12.26	20.39	18.14	12.47	10.32	10.67	8.68	14.11		
	HEMBA1001635	3.29	3.32	8.82	2.2	2.68	3.13	2.05	0.94	1.44		

Table 358

	HENBA1001122	9.39	9.29	17.71	6.58	5.73	6.53	2.46	3.1	3.26	*	-
	HENBA1001123	17.13	17.37	24.47	15.29	11.78	13.71	12.33	9.75	10.61	*	-
5	HENBA1001133	8.92	6.97	12.93	3.69	4.11	4.57	3.05	2.9	3.3	*	-
	HENBA1001137	6.57	7.37	12.88	4.24	4.79	2.86	4.1	2.06	3.89	*	-
	HENBA1001140	9.64	11.4	16.66	10.87	12.26	10.82	8.45	8.07	5.83	*	-
	HENBA1001144	75.62	79.41	130.35	64.7	57.88	71.91	43.23	38.45	52.04	*	-
	HENBA1001145	228.88	419.26	297.93	34.88	83.53	106.81	28.23	79.02	93.39	*	-
10	HENBA1001158	11.25	10.45	12.32	5.71	7.87	10.11	6.5	4.55	5.45	**	-
	HENBA1001172	13.23	15.14	21.25	10.8	9.67	9.91	10.54	8.89	11.23	*	-
	HENBA1001174	4.77	4.96	13.93	3.36	4.72	5.71	5.53	3.53	3.57	*	-
	HENBA1001175	25.64	49.85	37.18	5.53	36.99	18.9	6.53	8.52	5.43	*	-
	HENBA1001182	253.75	199.19	217.61	213.83	179.06	196.11	166.5	128.3	113.47	*	-
	HENBA1001184	2.72	5.66	6.46	2.92	7.84	3.84	2.54	1.64	3.55	*	-
15	HENBA1001192	7.71	9.61	13.89	5.11	4.84	2.98	2.49	2.97	4.44	*	-
	HENBA1001196	20.51	33.03	45	22.81	18.19	24.13	7.47	12.08	16.7	*	-
	HENBA1001197	377.82	621.74	482.95	613.89	355.35	492.48	223.5	509.73	494.09	*	-
	HENBA1001208	6.77	4.91	9.09	4.11	4.25	4.69	3.47	2.44	4.54	*	-
	HENBA1001213	162.45	150.13	151.86	224.78	129.48	188.61	135.18	97.08	115.8	*	-
	HENBA1001214	10.4	15.18	18.46	6.51	7.45	6.56	4.99	3.77	6.34	*	-
20	HENBA1001221	12.82	12.25	13.21	2.1	6.95	5.45	2.31	0.75	2.8	**	-
	HENBA1001225	4.23	4.43	8.6	1.43	2.73	4.08	2.02	0.94	2.52	*	-
	HENBA1001226	43.15	51.42	64.65	33.28	48.87	38.83	29.67	18.32	24.38	*	-
	HENBA1001228	5.8	4.52	8.07	5.69	4.51	3.32	3.32	1.37	1.24	*	-
	HENBA1001229	171.43	141.71	227.65	215.79	153.09	225.88	105.81	85.9	109.68	*	-
	HENBA1001235	51.44	62.4	68.51	46.01	34.3	36.55	26.86	20.23	22.59	*	-
25	HENBA1001238	19.14	19.72	19.8	10.86	15.88	9.5	15.04	10.36	16.33	*	-
	HENBA1001242	153.09	161.43	160.67	152.69	145.07	151.85	99.19	75.76	86.89	**	-
	HENBA1001247	33.3	29.15	48.14	21.32	32.34	22.28	34.27	20.52	27.05	*	-
	HENBA1001253	159.45	91.8	142.61	116.02	138.28	127.87	95.67	74.41	74.86	*	-
	HENBA1001257	10.7	11.57	12.78	5.67	10.33	4.37	9.19	6.06	4.62	*	-
	HENBA1001261	6.83	4.18	8.42	7.74	2.14	4.76	3.41	1.78	5.45	*	-
	HENBA1001262	12	7.56	5.09	4.62	4.49	6.27	3.32	1.44	4.7	*	-
30	HENBA1001265	15.44	18.41	25	9.71	8.97	12.14	13.41	14.3	11.72	*	-
	HENBA1001266	22.75	26.99	30.66	16.84	17.03	12.35	18.86	10.28	14.84	*	-
	HENBA1001269	62.54	134.76	5.56	0.22	1.58	0.07	24.28	23.59	16.64	*	-
	HENBA1001272	5.33	4.56	13.33	2.87	3.94	3.83	3.1	1.67	1.95	*	-
	HENBA1001279	26.1	31.13	42.25	28.49	33.02	28.73	22.53	17.36	24.84	*	-
35	HENBA1001281	43.22	55.66	55.89	31.39	34.2	26.87	21.08	11.24	15.62	*	-
	HENBA1001286	42.89	33.78	59.69	41.46	36.64	48.02	50.11	34.66	41.6	*	-
	HENBA1001289	4.71	4.83	5.28	3.13	4.27	3.23	3.56	1.6	3.12	*	-
	HENBA1001291	36.57	24.72	38.55	19.37	18.57	13.13	19.39	15.96	11.69	*	-
	HENBA1001294	8.41	15.41	16.09	9.8	8.09	6.28	18.25	5.05	15.73	*	-
	HENBA1001296	7.97	20.06	12.99	5.33	6.56	3.52	5.19	2.03	4.63	*	-
40	HENBA1001297	10.62	19.22	19.19	4.41	7.4	5.61	6.49	1.83	3.17	*	-
	HENBA1001299	34.22	39.68	53.21	25.14	29.23	20.66	39.68	22.86	32.79	*	-
	HENBA1001302	25.32	40.78	38.96	17.94	26.15	22.11	18.04	11.91	7.95	*	-
	HENBA1001303	10.97	8.86	13.92	4.53	5.04	4.42	4.71	4.03	7.45	*	-
	HENBA1001306	14.82	11.94	17.26	5.51	7.45	6.38	3.83	5.28	4.41	**	-
	HENBA1001308	44.73	35.49	88.39	29.14	49.52	28	40.25	28.31	38	*	-
45	HENBA1001310	8.14	6.04	8.8	2.67	3.81	3.08	5.37	3.64	2.99	**	-
	HENBA1001312	42.22	39.35	44.87	10.33	40.14	18.42	38.81	18.56	13.38	*	-
	HENBA1001319	3.83	6.75	12.77	3	5.03	13.5	4.72	0.95	4.62	*	-
	HENBA1001322	10.16	10.91	17.38	3.57	7.15	4.09	5.93	2.59	3.36	*	-
	HENBA1001323	41.14	36.93	47.79	17.38	18.24	15.31	13.85	10.42	3.9	**	-
	HENBA1001326	38.26	32.16	36.43	4.89	17.62	29.01	7.49	3.64	9.8	**	-
50	HENBA1001327	22.04	9.71	15.11	7.86	12.59	16.77	4.84	4.12	2.75	*	-
	HENBA1001330	29.76	31.78	64.11	34.57	26.99	26.81	34.76	20.01	20.9	*	-
	HENBA1001348	8.77	6.35	12.87	5.62	7.19	5.23	5.41	4.95	2.78	*	-
	HENBA1001350	11.78	12.52	19.81	3.72	3.89	6.89	7.45	5.17	4.88	*	-
	HENBA1001351	31.73	33.01	83.64	16.41	14.04	17.67	14.45	12.88	6.51	*	-
	HENBA1001352	16.11	10.31	24.58	3.9	5.27	4.13	8.48	3.75	2.44	*	-
55	HENBA1001353	239.33	87.11	196.31	30.75	66.77	70.17	148.92	56.48	39.11	*	-
	HENBA1001358	74.35	56.7	93	21.23	34.67	61.64	20.12	9.88	14.9	**	-
	HENBA1001361	9.19	10.7	19.27	11.26	10.32	15.03	5.68	3.47	4.41	*	-

Table 357

	HEMBA1000843	12.55	15.69	27.26	12.49	12.36	9.92	12.67	6.01	13.01			
	HEMBA1000851	7.99	8.88	11.08	3.31	3.75	3.94	6.33	2.77	2.83	**	*	-
5	HEMBA1000852	16.57	13.81	- 23	7.9	10.22	6.68	11.13	8.47	9.44	*	*	-
	HEMBA1000867	6.41	40.09	21.59	7.82	12.17	12.73	16.16	8.86	5.82			
	HEMBA1000869	6.66	6.92	14.84	4.94	6.58	4.87	5.66	1.59	2.82			
	HEMBA1000870	16.63	14.8	15.74	6.74	7.25	8.9	6.35	3.09	4.06	**	**	-
	HEMBA1000872	19	16.2	19.95	12.25	11.2	8.26	7.42	4.77	5.4	**	**	-
10	HEMBA1000875	7.38	6.41	11.16	2.07	4.19	4.92	3.96	2.9	2.7		*	-
	HEMBA1000876	12.29	12.13	19.01	9.58	10.69	6.67	12.67	6.89	9.73			
	HEMBA1000907	4.99	12.04	11.83	5.83	7.88	10.61	8.24	2.82	6.62			
	HEMBA1000908	7.89	3.64	11.79	2.7	10.96	1.89	2.83	0.62	3.13			
	HEMBA1000910	4.91	8.05	12.14	3.81	5.91	4.56	6.84	3.7	4.74			
	HEMBA1000918	21.04	31.68	45	25.17	23.52	19.82	25.88	9.86	18.68			
15	HEMBA1000919	4.44	2.5	5.8	3.35	2.86	3.17	3.08	0.43	2.37			
	HEMBA1000934	4.04	4.57	5.7	3.99	3.21	2.45	2.55	1.54	2.64	*		-
	HEMBA1000935	4.4	5.61	6.39	3.11	4.92	2.1	3.77	2.49	2.75	*		-
	HEMBA1000940	8.69	8.39	12.11	6.45	5.94	4.28	6.33	4.58	4.22	*	*	-
	HEMBA1000942	12.05	15.91	20.05	9.92	11.92	8.31	14.27	9.3	8.99			
	HEMBA1000943	2.61	3.51	9.04	0.06	1.63	0.43	1.65	0.02	0.85			
20	HEMBA1000946	5.48	9.79	16.74	3.82	4.51	6.06	3.38	0.73	4.11			
	HEMBA1000960	30.15	53.34	82.32	44.07	41.58	37.38	55.19	30.36	32.98			
	HEMBA1000962	10.91	13.68	14.99	4.87	9.87	8.15	6	5.38	7.85	*	**	-
	HEMBA1000968	9.18	5.88	12.53	7.34	5.67	5.5	4.92	3.16	5.55			
	HEMBA1000971	12.8	9.28	22.24	4.73	7.24	3.99	4.56	3.74	2.56	*		-
	HEMBA1000972	8.17	5.02	15.62	6.78	5.58	5.96	5.46	4.31	5.77			
25	HEMBA1000974	17.96	18.06	33.14	22.9	19.86	18.01	13.64	8.08	13.41			
	HEMBA1000975	8.54	6.18	15.36	5.78	4.48	4.06	6.6	5	4.55			
	HEMBA1000979	12.05	14.89	32.47	11.86	10.02	10.56	16.5	7.76	11.37			
	HEMBA1000981	31.09	24.35	60.14	12.11	16.96	23.49	26.22	14.59	14.47			
	HEMBA1000983	16.01	16.07	24.38	8.83	12.27	20.3	15.26	9.78	16.85			
	HEMBA1000985	7.28	3.65	6.98	4.91	4.64	4.99	4.05	3.11	1.91			
30	HEMBA1000986	13.51	10.48	26.88	15.8	12.5	15.39	12.22	6.19	7.06			
	HEMBA1000991	15.73	13.96	24.7	13.38	13.44	8.96	12.92	7.37	7.9			
	HEMBA1001007	6.2	4.4	10.1	3.64	3.96	2.18	2.52	1.45	2.54			
	HEMBA1001008	5.26	4.15	10.22	2.95	3.18	4.11	4.27	1.48	2.86			
	HEMBA1001009	2.98	2.47	7.99	2.44	3.53	2.07	2.05	0.9	2.64			
	HEMBA1001014	36.78	43.69	57.32	33.05	32.39	34.29	37.95	15.24	32.56			
35	HEMBA1001017	7.14	5.71	12.9	9.32	5.72	6.57	4.21	2.97	4.79			
	HEMBA1001019	6.35	3.7	5.48	7.22	3.14	5.53	3.87	2.27	2.61			
	HEMBA1001020	9.13	7.41	10.99	8.19	4.18	5.07	8.01	5.07	4.27			
	HEMBA1001021	4.2	4.32	9.05	3.87	3.59	5.86	6.3	1.95	6.15			
	HEMBA1001022	10.5	4.41	10.29	9.91	9.22	14.59	10.15	4.7	6.8			
	HEMBA1001024	1.82	2.83	5.27	3.68	2.95	3.77	2.92	1.3	2.29			
40	HEMBA1001026	3.74	3.71	5.2	2.89	4.11	4.69	3.01	1.9	3.66			
	HEMBA1001043	6.37	5.28	6.61	4.42	5.68	7.08	4.73	0.86	3.01			
	HEMBA1001051	120.57	138.21	211.05	111.88	82.79	99.92	82.72	88.42	94.95			
	HEMBA1001052	6.69	6.44	5.5	6.15	5.41	3.95	3.07	3.55	4.05	**		-
	HEMBA1001059	56.7	56.44	76.92	96.6	52.77	73	38.68	40.32	41.94	*		-
45	HEMBA1001060	22.93	14.44	26.88	16.88	16.28	14.69	14.87	11.27	12.22			
	HEMBA1001064	7.94	5.9	11.63	5.61	6.84	7.38	5.01	1.85	2.88			
	HEMBA1001071	2.87	5.22	8.35	8.63	6.39	5.39	3.26	1.85	3.81			
	HEMBA1001077	6.95	10.47	5.65	10.99	11.81	13.69	3.82	3.2	5.87			
	HEMBA1001078	13.55	16.29	18.17	15.7	24.22	16.62	14.25	7.88	7.1			
	HEMBA1001080	18.26	14.99	25.65	14.28	9.2	7.86	14.01	11.12	10.13			
	HEMBA1001084	10.12	12.02	19.64	11.43	8.23	10.4	8.52	5.91	8.47			
50	HEMBA1001085	12.67	14.13	17.46	10.37	7.16	7.36	9.72	8.19	7.2	*	*	-
	HEMBA1001088	3.47	2.77	6.21	3.04	2.22	2.64	4.71	4.47	2.67			
	HEMBA1001093	5.99	7.48	12.08	5.52	4.12	4.26	4.75	3.73	5.78			
	HEMBA1001094	3.87	3.67	7.3	1.83	2.98	1.32	1.79	1.61	3.04			
	HEMBA1001099	4.27	4.88	10.87	6.66	5.88	4.18	3.44	1.48	3.72			
	HEMBA1001104	3.23	5.58	9.6	7.66	21.55	10.79	5.83	1.68	4.13			
55	HEMBA1001109	55.09	49.61	96.04	54.26	41.61	45.47	44.91	34.62	57.09			
	HEMBA1001114	101.03	160.97	118.61	64.42	78.99	99.5	43.15	105.03	85.05			
	HEMBA1001121	8.11	7.04	9.73	5.44	4.04	4.32	4.43	2.6	4.72	*	*	-

Table 356

	HENBA1000551	31.43	29.13	74.52	42.08	38.34	36.48	27.97	18.18	17.63			
	HENBA1000555	23.07	51.65	28.63	16.75	19.18	84.47	21.47	21.24	16.53			
5	HENBA1000567	20.2	22.07	26.42	16.84	15.85	18.08	21.6	11.99	15.53	*	-	
	HENBA1000561	6.79	6.3	13.7	3.18	1.89	4.15	4.27	2.09	3.68			
	HENBA1000563	5.71	6.02	9.65	3.7	5.59	3.42	4.89	1.86	4.27			
	HENBA1000567	2.91	1.55	3.34	2.35	0.99	3.67	2.89	1.41	2.17			
	HENBA1000568	8.19	5.62	14.84	8.57	7.03	7.81	7.88	3.61	9.31			
10	HENBA1000569	2.57	3.31	7.08	4.64	2.77	3.17	2.3	1.18	2.24			
	HENBA1000575	24.44	15.32	39.5	24.17	18.92	19.53	17.36	14	17.06			
	HENBA1000588	7.1	5.81	8.91	7.39	8.88	8.85	8.33	2.25	5.9			
	HENBA1000590	5.23	6.06	5.7	5.36	6.93	6.66	4.07	2.31	4.39	*	-	
	HENBA1000591	24.12	22.46	26.56	10.74	18.87	21.61	11.03	5.76	13.01	**	-	
	HENBA1000592	30.95	41.43	24.53	10.67	31.41	23.82	10.57	11.36	12.37	*	-	
15	HENBA1000594	5.11	7.29	6.57	8.37	4.71	6.68	5.32	3.27	3.89			
	HENBA1000604	21.62	17.48	22.7	28.09	18.45	20.99	13.92	9.43	17.25			
	HENBA1000607	59.03	59	91.06	71.67	51.82	56.83	60.28	42.6	51.42			
	HENBA1000608	9.41	4.42	6.41	8.42	5.12	4.07	7.92	3.99	8.94			
	HENBA1000622	7.92	7.76	16.15	9.11	9.27	13.89	9.94	3.81	11.14			
	HENBA1000634	52.78	74.71	70.68	39.88	65.82	64.7	32.15	24.17	25.34	**	-	
20	HENBA1000636	27.22	28.11	25.27	25.4	22.96	20.49	9.12	9.28	12.36	**	-	
	HENBA1000637	8.86	11.75	7.97	5.21	9.26	6.88	4.93	3.34	3.68	*	-	
	HENBA1000655	30.73	31.69	54.9	27.54	16.27	22.75	21.5	18.39	24.41			
	HENBA1000657	14.08	9.28	10.28	5.96	7.53	6.3	3.93	18.82	5.24	*	-	
	HENBA1000662	6.27	5.79	10.27	3.02	6.27	5.29	2.93	4.16	3.38			
	HENBA1000664	6.29	6.77	9.71	5.57	5.25	9.62	2.22	2.8	4.62	*	-	
25	HENBA1000671	30.41	19.43	27.86	31.58	22.01	24.63	20.58	14.98	15.56			
	HENBA1000673	24.11	24.74	28.83	18.72	18.93	18.52	21.96	17.14	28.31	**	-	
	HENBA1000675	11.22	13.93	16.24	11.01	14.94	12.36	11.46	6.91	7.33			
	HENBA1000678	5.24	8.15	11.3	5.35	7.05	6.17	5.02	1.55	1.87			
	HENBA1000682	34.57	35.39	53.17	14.17	19.8	19.96	13.73	23.76	33.18	*	-	
	HENBA1000686	21.71	40.16	35.85	10.91	16.11	14.54	4.3	11.52	5.8	*	*	-
30	HENBA1000702	10.84	17.35	25.73	7.39	11.25	13.13	7.56	7.08	9.42			
	HENBA1000705	4.6	3.75	8.97	3.21	1.09	3.11	3.08	4.72	3.61			
	HENBA1000713	10.43	11	19.66	5.79	9.41	8.35	11.72	6.95	13.37			
	HENBA1000718	27.65	31.72	34.47	18.34	17.98	21.82	22.89	15.55	26.61	**	-	
	HENBA1000719	21.36	28.6	18.8	18.63	23.29	31.74	21.77	16.83	13.89			
	HENBA1000722	10.72	14.07	15.25	12.4	11.9	8.36	7.24	4.41	3.46	**	-	
35	HENBA1000726	32.15	32.06	69.89	34.36	26.18	30.36	36.32	22.04	41.04			
	HENBA1000727	16.97	26.82	19.68	5.88	11.41	11.31	4.45	8.11	7.29	*	**	-
	HENBA1000732	8.15	9.73	16.83	6.74	7.08	9.59	5.57	4.8	5.38			
	HENBA1000736	9.38	7.92	11.45	7.38	12.77	8.49	9.33	6.94	12.57			
	HENBA1000743	6.76	5.78	11.72	5.12	7.77	27.92	5.16	3.39	8.47			
	HENBA1000745	7.61	8.54	13.88	4.04	5.82	4.46	4.98	2.87	5.45			
40	HENBA1000747	5.32	4.72	12.62	2.69	5.68	3.35	3.85	2.1	3.4			
	HENBA1000748	7.08	6.97	13.44	6.2	9.91	9.41	8.23	3.96	6.55			
	HENBA1000749	18.03	20.17	33.28	13.55	12.82	15.47	12.43	12.59	18.76			
	HENBA1000752	9.61	11.01	10.43	7	5.98	6.12	4.8	4.54	8.58	**	*	-
	HENBA1000753	14.73	8.44	14.96	4.78	5.04	11.61	5.39	6.35	8.17			
	HENBA1000757	11.82	6.81	16.04	4.91	5.14	4.63	2.56	3.42	2.84	*	-	
45	HENBA1000760	13.51	13.09	20.98	7.09	10.48	10.58	5.68	4.5	6.77	*	-	
	HENBA1000769	7.76	13.89	13.94	4.47	6.26	7.04	9.68	6.14	7.78			
	HENBA1000773	3.65	4.13	15.13	0.99	5.68	8.11	2.71	1.47	3.38			
	HENBA1000774	21.34	27.53	42.71	22.13	28.3	22.38	30.47	15.88	17.88			
	HENBA1000780	6.15	5.35	9.22	6.03	4.9	4.38	2.5	2.74	2.78	*	-	
	HENBA1000783	8.54	7.55	8.23	4.94	5.39	5.68	3.76	3.66	3.46	**	**	-
50	HENBA1000791	29.92	28.02	44.77	28.48	22.57	21.5	32.85	19.21	32.35			
	HENBA1000793	30.39	19.19	19.38	8.64	16.27	16.1	13.32	16.07	12.64			
	HENBA1000802	4.74	2.72	8.7	3.43	2.84	2.21	2.09	1.11	2.74			
	HENBA1000813	46.62	47.69	50.56	27.4	35.01	30.28	30.89	15.81	34.56	**	*	-
	HENBA1000817	12.55	12.98	16.93	6.72	11.22	3.95	2.79	1.89	3.94	**	-	
	HENBA1000822	6.54	19.54	13.74	4.86	7.54	6.86	4.29	1.48	3.36			
55	HENBA1000827	13.88	14.84	28.11	13.31	20.08	16.37	7.68	5.83	12.73			
	HENBA1000833	12.7	15.8	16.64	5.65	5.56	4.66	3.19	1.79	1.87	**	**	-
	HENBA1000835	118.36	117.03	136.17	175.69	117.43	151.33	112.03	74.79	83.7			

Table 355

	HENBA1000290	5.54	5.24	10.06	5.33	1.73	3.05	3.54	2.47	2.22			
	HENBA1000296	9.5	13.84	20.76	5.15	12.54	6.86	5.85	5.2	3.4	*	-	
5	HENBA1000300	35.72	34.39	45.53	27	31.39	25.7	26.91	18.61	28.62	*	-	
	HENBA1000302	3.92	5.7	9.9	1.71	5.14	2.38	3.51	1.14	3.24			
	HENBA1000303	7.88	7.6	13.76	3.6	4.76	4.02	4.64	2.88	3.92	*	-	
	HENBA1000304	20.77	16.02	19.73	12.6	11.48	11.65	12.2	13.31	13.14	**	*	-
	HENBA1000307	7.67	8.03	6.72	5.49	4.16	4.06	1.61	1.51	3.67	**	**	-
10	HENBA1000312	26.83	27.01	36.09	21.32	17.54	18.68	18.71	13.82	23.54	*	-	
	HENBA1000318	14.81	10.63	16.92	6.25	7.77	8.66	2.82	1.83	3.66	*	**	-
	HENBA1000327	16.1	19.56	31.73	13.19	19.56	35.1	16.48	11.24	13.77			
	HENBA1000333	4.8	4.82	8.3	2.23	3.1	2.42	2.41	0.71	1.87	*	*	-
	HENBA1000338	14.85	18.8	25.1	10	13.67	12.74	13.23	10.79	10.59			
	HENBA1000343	9.76	12.6	14.72	7.21	10.87	8.74	6.58	5.91	8.76	*	-	
15	HENBA1000349	7.48	9.48	16.3	4.37	4.58	5.11	3.79	2.9	11.15			
	HENBA1000351	16.95	12.69	26.79	18.1	17.55	14.44	11.22	7.92	17.2			
	HENBA1000355	11.99	7.36	8.03	7.48	3.22	2.44	1.27	2.39	3.56	*	-	
	HENBA1000356	34.47	36.59	32.24	39.62	29.15	32.82	22.27	24.67	20.29	**	-	
	HENBA1000357	16.16	16.75	30.37	11.91	12.99	11.46	14.62	9.28	12.57			
	HENBA1000366	10.73	7.46	15.42	5.17	5.07	3.55	5.42	3.22	6.53	*	-	
20	HENBA1000369	10.72	11.69	16.34	4.55	5.5	4.39	6.76	3.13	5.67	*	*	-
	HENBA1000370	9	13.55	13.78	4.72	8.24	7.37	5.92	3.13	3.05	*	*	-
	HENBA1000376	90.99	73.26	0.51	1.99	5.8	2.22	56.72	52.09	77.09			
	HENBA1000387	26.37	30.02	2.06	0.67	28.79	0	22.74	18.21	26.75			
	HENBA1000389	16.44	15.64	23.44	8.78	10.05	15.24	16.41	8.42	9.96			
	HENBA1000390	10	6.3	9.6	4.59	5.15	2.55	7.47	3.11	6.96	*	-	
25	HENBA1000392	7.7	8.4	14.19	5.19	4.42	2.87	4.91	3.87	4.75			
	HENBA1000396	9.8	6.67	9.18	1.73	1.25	0.34	3.88	1.99	4.11	**	*	-
	HENBA1000411	4.32	4.84	13.72	2.13	3.17	1.91	3.36	1.45	2.17			
	HENBA1000418	10.46	14.65	17.48	5.45	8.2	5.71	4.56	2.35	3.71	*	**	-
	HENBA1000422	9.96	6.26	0	0.69	0.58	0.28	3.62	3.45	7.58			
	HENBA1000428	8.49	7.69	1.25	0.25	8.33	0	5.22	5.3	9.15			
30	HENBA1000434	6.78	3.73	4.63	2.97	2.1	1.84	5.95	1.48	3.75	*	-	
	HENBA1000442	3.73	3.34	8.65	2.11	3.23	1.05	2.87	2.62	6.58			
	HENBA1000443	6.03	7.17	14.31	3.08	6.26	3.05	8.49	4.27	7.22			
	HENBA1000446	56.88	47.2	7.47	1.38	1.35	0.83	46.56	24.3	38.67			
	HENBA1000456	14.88	20.85	28.05	5.95	15.86	10.1	10.12	2.72	7.69	*	-	
	HENBA1000459	15.34	18.32	27.74	11	17.98	14.9	11.74	7.37	11.06			
35	HENBA1000460	23.55	16.11	28.93	19.24	20.63	19.37	10.12	7.72	17.35			
	HENBA1000462	25.82	7.48	2.26	1.21	3.05	10.25	13.12	8.17	7.37			
	HENBA1000464	7.56	4.51	6.06	3.68	3.64	2.38	3.55	1.26	2.53	*	*	-
	HENBA1000468	3	4	6.33	1.85	2.29	0.39	1.45	0.17	1.64	*	*	-
	HENBA1000469	22.54	28.03	34.83	17.42	17.27	14.02	27.65	14.61	18.23	*	-	
	HENBA1000477	17.38	15.06	7.59	1.07	12.5	0.35	11.11	7.11	9.19			
40	HENBA1000481	63.47	80.18	49.05	5.51	28	7.01	17.9	5.67	5.55	*	**	-
	HENBA1000488	14.11	13.36	8.09	0.4	9.3	4.86	6.34	1.81	3.54	*	*	-
	HENBA1000490	9	6.13	11.44	3.85	4.12	5.04	6.84	2.63	7.56	*	-	
	HENBA1000491	5.32	6.61	17.42	8.34	8.06	9.74	6.93	4.28	9.22			
	HENBA1000498	14.5	18.25	24.06	10.95	15.11	14.4	16.82	10.93	14.38			
	HENBA1000501	7.87	9.11	2.74	0.21	2.74	4.51	6.04	4.62	6.97			
45	HENBA1000504	12.54	5.58	11.75	3.41	4.13	2.59	2.67	4.18	4.58	*	-	
	HENBA1000505	6.92	8.54	7.08	0	1.28	0.84	5.6	2.13	4.7	**	*	-
	HENBA1000507	0.45	1.56	8.28	0	1.2	0.18	0.77	0	0.38			
	HENBA1000508	26.91	34.97	50.59	22.03	32.11	27.29	38.18	21.05	20.98			
	HENBA1000518	3.76	3.28	6.48	5.32	4.44	6.01	3.59	1.86	3.38			
	HENBA1000519	85.07	91.86	141.09	42.18	50.11	71.14	47.48	35.9	57.59	*	-	
50	HENBA1000520	6.49	7.75	13.06	6.32	6.26	4.22	5.77	4.5	4.14			
	HENBA1000523	11.85	6.14	1.44	0	1.42	0.29	4.15	3.58	2.35			
	HENBA1000531	8.26	6.64	16	4.73	9.37	11.46	9.54	4.57	6.14			
	HENBA1000534	28.94	37.16	55.86	24.81	22.76	20.48	41.37	29.08	32.84			
	HENBA1000538	14.97	19.18	30.49	12.11	14.95	11.04	13.25	9.03	10.27			
	HENBA1000540	18.41	10.82	26.42	7.05	8.02	12.43	11.49	5.58	4.88			
55	HENBA1000542	68.21	65.66	96.9	105.13	126.6	107.82	49.4	38.5	50.6	*	*	+
	HENBA1000545	10.72	9.72	19.53	13.15	9.91	10.84	7.59	4.75	4.2			
	HENBA1000547	16.09	12.61	26.19	11.54	11.54	13.04	5.83	5.77	5.14	*	-	

Table 354

	BRAWH1000096	5.38	6.14	9.95	5.23	5.99	6.41	5.29	2.17	4.29			
	BRAWH1000097	32.73	36.25	38.91	16.1	26.45	25.46	34.97	22.26	21.98	*	-	
5	BRAWH1000100	3.44	3.77	5.35	7.33	1.79	1.51	2.06	0.88	1.83	*	*	-
	BRAWH1000101	151.17	188.03	182.43	124.31	137.07	112.48	156.93	114.91	72.6	*	-	
	BRAWH1000104	3.11	5.16	13	2.27	6.18	12.04	5.42	9.52	5.16			
	BRAWH1000107	4.13	9.11	10.29	5.8	8.93	4.46	6.91	1.75	1.45			
	BRAWH1000110	63.35	48.25	85.05	53.28	34.71	42.03	42.56	51.83	49.06			
	BRAWH1000111	17.89	14.24	19.42	6.38	13.57	14.32	11.88	11.36	7.34	*	*	-
10	BRAWH1000135	7.97	7.93	10.18	6.35	4.56	3.75	5.18	4.92	2.5	*	*	-
	BRAWH1000190	13.15	11.52	18.33	11.13	9.11	7.4	12.15	9.59	8.37			
	HEMBA1000005	35.95	34.82	38.35	20.64	26.11	18.05	30.81	17.47	24.15	**	*	-
	HEMBA1000006	9.78	9.8	11.36	1.4	5.53	2.46	3.8	1.71	7.12	**	*	-
	HEMBA1000012	142.4	148.44	163.52	136.28	167.18	112.45	92.75	76.02	85.25	**		-
15	HEMBA1000020	190.7	248.05	252	200.63	259.1	183.87	186.01	114.55	161.9			
	HEMBA1000030	20.38	15.29	27.68	14.24	16.3	11.52	9.94	9.08	14.21			
	HEMBA1000034	15.22	17.47	15.15	7.36	10.81	14.4	7.69	4.87	8.91	**		-
	HEMBA1000042	22.23	17.73	35.18	17.22	20.39	12.68	19.66	13.46	21.05			
	HEMBA1000045	12.71	14.02	2.46	0.7	7.88	1.39	4.27	2.85	2.17			
	HEMBA1000046	10.19	13.7	20.59	6.92	10.64	7.67	13.72	12.92	11.87			
20	HEMBA1000047	4.05	7.12	13.48	2.54	4.15	2.97	2.37	0.81	3.55			
	HEMBA1000048	10.63	14.86	15.51	2.18	6.64	3.23	4.43	0.94	3.67	**	**	-
	HEMBA1000050	3.15	3.21	10.5	2.13	3.59	2.35	2.36	0.05	2.24			
	HEMBA1000053	5.98	3.98	9.78	8.93	5.91	6.48	2.35	1.85	5.68			
	HEMBA1000060	10.62	11.2	18.18	10.13	8.71	11.8	7.34	2.99	9.16			
	HEMBA1000072	530.32	492.25	767.01	366.98	521.7	354.83	288.04	399.85	194.47	*		-
25	HEMBA1000073	26.51	17.12	39.76	19.17	28.84	19.97	20.8	18.82	12.3			
	HEMBA1000076	30.89	23.42	46.09	20.74	16.14	16.31	22.31	20.51	20.54			
	HEMBA1000084	46.03	48.38	54.47	61.64	67.48	73.34	43.23	26.25	43.33	*	+	
	HEMBA1000087	2.35	3.13	9.77	1.66	2.08	0.86	1.69	0.8	1.37			
	HEMBA1000088	3.05	3.73	12.72	2.97	3.67	2.66	4.68	0.94	2.56			
	HEMBA1000091	38.82	23.63	75.02	58.3	74.93	71.96	24.1	16.38	30.08			
30	HEMBA1000111	21.39	20.11	44.78	24.89	20.7	26.97	20.62	12.93	18.49			
	HEMBA1000121	18.15	14.06	45.72	26.38	18.11	19.11	16.09	9.88	11.04			
	HEMBA1000128	6.74	4.73	14.38	8.16	5.93	6.25	3.31	1.93	2.79			
	HEMBA1000129	6.94	5.68	13.44	7.51	6.89	6.55	4.13	5.51	3.96			
	HEMBA1000141	12.4	7.44	22.8	8.83	6.54	5.23	5.19	3.51	3.66			
	HEMBA1000146	6.55	7.84	15.71	4.62	7.62	7.46	5.3	2.57	3.09			
35	HEMBA1000150	28.41	41.35	48.05	22.83	16.47	23.82	28.66	15.99	24.53	*		-
	HEMBA1000154	717.61	696.56	881.96	454.74	531.19	670.47	527.79	542.09	612.33	*	*	-
	HEMBA1000156	10.8	18.5	21	9.98	4.23	6.43	3.69	3.2	3.36	*	*	-
	HEMBA1000158	18.74	13.45	14.16	7.7	2.81	4.64	5.39	2.06	3.26	**	**	-
	HEMBA1000168	8.01	6.75	7.08	5.88	4.99	5.86	6.83	2.42	4.44	*		-
	HEMBA1000180	3.02	3.26	4.77	4.2	2.93	2.73	2.69	1.65	3.39			
40	HEMBA1000185	19.4	18.46	23.05	27.72	15.45	13.55	13.51	10.97	19.3			
	HEMBA1000188	6.28	5.61	7.48	5.62	6.91	3.78	3.34	2.85	4.56	*		-
	HEMBA1000193	7.66	7.79	11.97	13.74	5.68	7.85	5.99	5.1	6.65			
	HEMBA1000194	31.21	25.82	29.1	20.52	12.15	13.24	9	11.15	12.72	*	**	-
	HEMBA1000201	13.81	8.63	14.31	11.15	8.36	7.64	5.43	4.05	4.24	*		-
	HEMBA1000213	2.97	4.59	8.02	5.36	2.78	4.45	3.2	1.24	3.2			
45	HEMBA1000216	7.62	8.45	11.73	6.7	6.96	6.62	6.13	4.35	5.56	*		-
	HEMBA1000227	2.73	4.56	8.5	4.18	3.58	2.84	3.35	1.71	2.27			
	HEMBA1000231	24.52	27.31	24.37	15.72	17.43	21.38	21.29	12.5	20.49	*		-
	HEMBA1000237	76.81	74.32	83.86	66.84	69.97	42.93	44.61	22.46	38.37	**		-
	HEMBA1000243	13.82	10.47	25.88	21.77	11.3	17.41	12.09	8.84	11.38			
	HEMBA1000244	9	7.04	9.25	7.59	2.86	4.81	3.47	4.91	3.18	**		-
50	HEMBA1000251	4.62	4.41	5.87	4.22	1.63	3.25	2.37	2.22	2.32	**		-
	HEMBA1000254	10.09	13.73	13.57	7.22	5.82	7.35	8.91	6.98	9.5	*	*	-
	HEMBA1000264	5.1	3.66	11.16	4.22	2.15	2.47	2.98	1.73	5.35			
	HEMBA1000269	4.34	6.24	5.63	3.94	4.73	5.35	3.13	1.23	3.19	*		-
	HEMBA1000275	22.4	40.89	22.83	21.15	22.57	23.38	10.3	5.45	12.41	*		-
	HEMBA1000280	6.51	8.26	15.12	7.28	7.37	8.84	7.59	2.9	7.78			
55	HEMBA1000282	18.71	16.98	38.85	19.86	12.71	13.67	18.74	14.02	21.9			
	HEMBA1000287	10.5	12.84	12.5	3.76	7.11	6.81	3.4	8.14	8.93	**		-
	HEMBA1000288	14.15	21.12	36.73	13.92	11.11	18.3	14.48	11.79	19.51			

	Clone	UV 0h			UV 4h			UV 24h			t test		4h 24h
		Exp. 1	Exp. 2	Exp. 3	Exp. 1	Exp. 2	Exp. 3	Exp. 1	Exp. 2	Exp. 3	0/4	0/24	+/- +/-
5	GAPDH(Cr1)	0	1.29	0.1	0.9	0.06	1.18	1.49	0.47	0			
	β actin(Cr2)	256.82	283.53	414.29	388.38	117.29	329.8	189.18	190.26	157.87	*		-
	ADRGL1000005	15.9	10.68	19.67	3.3	6.44	12.69	6	3.06	5	*		-
	ADRGL1000007	47.47	31.62	85.7	41.54	41.35	40.62	31.64	14.77	31.5			
	ADRGL1000009	3.97	2.45	8.72	6.06	2.65	3.32	3.72	1.8	2.91			
10	ADRGL1000011	21.55	13.26	21.8	18.36	12.63	13.95	15.62	6.5	14.45			
	ADRGL1000027	2.4	1.68	5.6	4.02	2.48	2.04	3.18	0.81	3.55			
	ADRGL1000058	84.24	29.04	76.55	95.27	68.98	72.58	39.84	22.31	39.98			
	ADRGL1000069	7.61	6.31	10.67	3.58	5.86	4.59	2.92	1.14	2.62	*		-
	ADRGL1000077	4.08	5.99	6.34	5.76	3.87	4.68	4.23	2	3.08			
	ADRGL1000092	41.45	38.57	55.77	73.06	40.91	54.59	24.67	18.9	27.36	*		-
15	ADRGL1000099	43.57	33.01	48.6	56.04	29.24	45.47	24.87	18.08	20.27	*		-
	ADRGL1000136	52.22	43.6	70.27	58.84	40.95	44.91	35.09	21.07	29.76	*		-
	ADRGL1000147	6.66	6.78	11.64	6.52	5.38	7	5.38	2.59	5.89			
	ADRGL1000159	7.28	7.31	7.55	10.23	10.38	9.89	7.3	4.68	8.62	**		+
	ADRGL1000160	11.62	12.89	8.28	9.39	13.77	23.6	7.39	5.18	9.59			
20	ADRGL1000171	4.47	4.72	5.01	4.83	6.43	4.19	3.69	1.77	4.29			
	ADRGL1000181	9.73	12.6	18.02	18.65	11.48	13.93	9.96	11.18	7.77			
	BGG111000015	25.13	21.03	22.91	10.68	12.73	20.09	10.32	15.5	18.5	*		-
	BGG111000016	63.98	55.25	74.82	18.05	36.94	8.26	17.67	40.65	37.24	*	*	-
	BGG111000017	8.13	6.51	11.46	3.65	5.27	5.54	2.72	4.69	3.8	*		-
	BGG111000022	12.98	14.45	15.11	7.84	9.62	8.77	7.97	8.92	10.08	**	**	-
25	BGG111000031	13.67	9.32	14.45	9.12	13.53	12.58	6.75	10.28	15.5			
	BGG111000042	78.81	92.28	77.71	51.19	53.47	19.5	23.41	16.53	22.63	*	**	-
	BGG111000046	59.14	59.17	43.06	30.88	42.51	35.11	16	15.29	7.23	**		-
	BNGH41000025	34.23	58.36	71.87	11.96	22.6	19.45	12.51	15.56	19.08	*	*	-
	BNGH41000026	8.61	10.35	11.92	5.79	4.65	4.41	2.7	3.65	4.11	**	**	-
30	BNGH41000027	36.04	26.15	57.91	21.21	34.28	46.33	28.86	28.53	19.37			
	BNGH41000035	71.93	95.4	103.3	77.48	91.38	82.2	85.4	91.4	89.21			
	BNGH41000037	11.37	8.43	18.43	4.88	12.04	10.79	4.26	3.88	5.78			
	BNGH41000042	153.34	222.69	94.88	128.41	120.85	102.56	66.52	26.75	52.39	*		-
	BNGH41000048	115.13	80.94	131.08	158.03	99.22	95.02	70.52	58.67	62.66	*		-
	BNGH41000056	6.81	7.33	32.46	17.81	11.49	11.22	6.84	4.36	6.34			
35	BNGH41000087	7.57	4.78	6.77	6.26	6.74	6.12	2.23	3.11	5.45			
	BNGH41000091	5.88	4.66	7.21	3.92	3.81	2.29	2.51	1.95	5.23	*		-
	BNGH41000157	24.78	9.57	39.68	15.95	28.36	14.05	14.96	12.43	25.4			
	BNGH41000169	4.77	2.7	8.76	3.03	4.07	2.47	2.48	1.85	2.4			
	BNGH41000181	15.03	14.16	15.82	9.14	8.43	9.02	9.57	5.48	4.7	**	**	-
	BNGH41000198	5.23	7.17	13.44	2.81	5.92	4	2.64	2.63	2.78			
40	BNGH41000219	55.36	63.96	42.98	30.63	34.86	27.19	11.04	4.75	5.11	*	**	-
	BNGH41000229	41.48	41.07	32.45	12.86	20.86	15.06	12.11	29.5	42.2	**		-
	BNGH41000237	30.57	28.92	29.88	23.95	17.14	16.36	7.62	11.39	18.2	*	**	-
	BNGH41000238	12.97	6.92	11.13	3.1	5.96	5.05	3.21	2.99	4.83	*	*	-
	BNGH41000243	37.29	22.23	38.12	21.69	20.76	22.29	17.48	13.04	12	*		-
45	BNGH41000270	7.24	2.74	13.66	5.03	4.81	3.91	2.55	1.06	3.15			
	BRAH1000004	26.05	12.1	22.36	14.77	13.7	15.56	10.21	9.28	9.3			
	BRAH1000018	16.02	18.04	23.1	10.35	11.87	13.62	8.4	6.81	8.26	*	**	-
	BRAH1000021	13.77	12.07	18.61	14.58	14.48	10.65	13.45	8.59	11.48			
	BRAH1000027	4.8	4.82	5.71	4.63	4.59	5.52	2.19	2.04	7.42			
	BRAH1000029	7.2	4.85	5.05	4.67	2.83	2.65	2.93	2.18	1.97	*		-
50	BRAH1000040	20.85	33.58	27.15	16.04	13.44	12.14	8.98	5.99	6.77	*	**	-
	BRAH1000050	86.78	63.26	107.91	121.47	88.32	92.39	77.36	45.55	64.9			
	BRAH1000051	3.25	3.15	8.62	2.34	1.27	1.81	2.29	1.14	1.52			
	BRAH1000060	103.56	87.24	102.14	122.05	99.97	107.12	58.34	59.07	66.98	**		-
	BRAH1000075	6.97	6.63	15	3.29	4.01	3.69	3.52	2.95	3.34			
	BRAH1000081	23.41	12.71	28.79	20.72	20.86	22.65	19.13	9.85	15.04			
55	BRAH1000084	219.94	140.27	3.27	8.21	5.55	1.75	106.89	121.2	155.45			
	BRAH1000095	6.67	5.04	6.84	7.47	5.22	4.62	3.81	0.91	1.79	*		-

Table 353

5 Expression of each cDNA in skin-derived fibroblast cells exposed and unexposed to
ultraviolet light (the table also includes clones that are not described in Examples)

10 In the table, UV_0h represents skin-derived fibroblast cells without ultraviolet irradiation;
UV_4h and UV_24h represent skin-derived fibroblast cells 4 and 24 hours, respectively, after the
irradiation. The assay was performed in triplicate (n=3) and each result is shown in the column
of Exp1, Exp2, or Exp3. "t-test 0/4" and "t-test 0/24" represent the results of the test for
significant difference between the unexposed cells and the cells 4 hours after the irradiation, and
15 between the unexposed cells and the cells 24 hours after the irradiation, respectively. The table
also includes the information on an increase (+) or decrease (-) in the expression level of the
gene in the exposed cells 4 hours or 24 hours after the ultraviolet light irradiation. The results of
the test for significant difference are shown in the columns of *p<0.05 and **p<0.01.
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	Y79AA1002365	0.66	2.04	2.26	1.97	3.51	2.25		
	Y79AA1002373	1.17	3.93	2.42	1.59	1.97	1.43		
5	Y79AA1002376	110.81	135.82	-249.8	205.99	213.25	191.69		
	Y79AA1002378	1.9	4.8	4.91	2.2	3.6	3		
	Y79AA1002381	8.65	14.11	19.19	18.84	21.52	17.97		
10	Y79AA1002388	7.05	9.99	18.24	15.88	21.51	19.99		
	Y79AA1002399	1.79	4.25	3.74	4.62	4.08	3.47		
	Y79AA1002407	3.05	4.16	3.13	4.66	5.77	4.5	*	+
15	Y79AA1002413	3.21	6.78	8.05	6.46	8.32	6.87		
	Y79AA1002416	1.46	5	2.74	2.49	3.44	3.55		
	Y79AA1002429	5.5	8.15	7.27	8	11.11	8.01		
20	Y79AA1002431	0.92	4.43	0.48	0.79	1.78	0.89		
	Y79AA1002433	1.27	5.9	3.24	4.8	3.84	5.58		
	Y79AA1002445	4.01	5.34	5.76	3.1	4.89	5.41		
25	Y79AA1002461	0.63	2.45	1.79	1.19	2.71	1.41		
	Y79AA1002466	39.02	70.71	94.5	91.12	82.27	94.71		
	Y79AA1002471	4.44	6.67	6.08	7.43	8.06	10.49		
	Y79AA1002472	2.41	6.16	5.99	6.8	8.39	4.06		
30	Y79AA1002474	1.93	8.27	4.31	4.89	6.52	7.13		
	Y79AA1002482	3.52	6.66	10.37	9.02	11.81	8.69		
	Y79AA1002487	1.38	4.12	2.46	1.96	3.01	2.56		
35	Y79AA1002490	10.37	9.91	16.35	11.11	12.88	16.86		
	Y79AA1002493	1.96	4.07	6.14	6.5	8.9	4.1		
	ZRV6C1006278	0.61	4.08	2.22	1.81	1.58	2.11		

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	Y79AA1002089	1.18	3.46	2.13	1.46	2.92	3.26		
	Y79AA1002093	2.19	5.48	5.25	5.28	5.68	6.17		
5	Y79AA1002101	1.11	8.58	-2.98	6.54	5.58	6.6		
	Y79AA1002103	1.47	10.22	3.39	4.43	6.49	4.7		
	Y79AA1002115	4.34	9.78	7.37	7.45	7.03	6.95		
	Y79AA1002121	1.55	2.16	2.18	1.67	2.55	2.31		
10	Y79AA1002125	6.67	7.08	9.29	8.81	6.4	7.6		
	Y79AA1002129	1.64	6.23	7.84	5.41	2.2	4.93		
	Y79AA1002131	0.9	3.25	0.78	0.77	1.29	1.15		
15	Y79AA1002139	0.69	5.02	1.04	1.83	1.53	1.34		
	Y79AA1002144	25.99	29.62	51.01	42.61	51.16	43.17		
	Y79AA1002177	1.72	5.97	4.33	4.79	3.09	4.73		
	Y79AA1002183	10.44	13.89	17.69	27.61	29.67	28.92	**	+
20	Y79AA1002202	3.97	7.15	8.34	18.27	10.12	17.85	*	+
	Y79AA1002204	0.53	0.99	1.56	1.7	2.2	1.54		
	Y79AA1002206	2.63	5.36	7.28	4.35	2.95	1.49		
	Y79AA1002208	4.26	6.54	3.94	6.88	6.3	3.96		
25	Y79AA1002209	1.8	6.34	2.88	4.38	3.74	4.57		
	Y79AA1002210	0.41	4.14	2.09	1.8	2.24	1.65		
	Y79AA1002211	2.25	5.39	3.85	5.71	5.3	4.5		
30	Y79AA1002213	1.15	4.13	6.53	7.38	7.54	7.43		
	Y79AA1002215	18.7	18.69	26.61	17.72	15.59	9.62		
	Y79AA1002220	3.78	3.38	2.87	4.89	4.19	4.14	*	+
	Y79AA1002226	8.54	8.9	9.75	13.06	14.2	4.41		
35	Y79AA1002229	1.35	3.88	3.38	2.95	2.79	2.67		
	Y79AA1002234	3.24	6.82	3.94	4.29	7.74	6.88		
	Y79AA1002235	5.6	7.55	6.43	8.78	9.74	9.47	*	+
40	Y79AA1002246	0.59	5.06	2.41	3.94	2.54	4.27		
	Y79AA1002258	0.72	7.26	2.92	3.99	4.19	2.7		
	Y79AA1002279	17.79	19.12	27.8	16.52	19.13	11.5		
	Y79AA1002292	1.68	2.1	3.22	2.96	3.91	2.73		
45	Y79AA1002298	0.76	2.52	1.32	2.03	2.77	1.06		
	Y79AA1002307	1.05	4.35	1.79	0.76	1.05	1.2		
	Y79AA1002309	1.15	4.19	2.3	2.21	1.78	2.55		
	Y79AA1002311	2.84	7.35	3.43	5.71	6.04	5.45		
50	Y79AA1002334	1.72	6.54	2.95	4.77	4.19	3.35		
	Y79AA1002351	1.27	5.5	2.89	3.5	3.38	3.06		
	Y79AA1002355	12.83	12.25	28.96	22.94	22.07	21.02		
55	Y79AA1002361	2.22	2.27	3.26	2.47	4.54	1.55		

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	Y79AA1001846	1.82	6.52	5.51	6.52	5.09	5.07		
	Y79AA1001848	0.86	5.57	2.75	5.98	4.78	5.14		
5	Y79AA1001853	1.38	6.16	2.76	3.24	4.56	3.63		
	Y79AA1001863	0.86	5	2.53	2.93	3.83	3.85		
	Y79AA1001866	2.29	3.2	5.81	3.53	3.98	3.46		
	Y79AA1001874	0.12	2.18	-0.21	1.17	0.75	0.51		
10	Y79AA1001875	9.33	12.67	13.09	11.05	17.79	18.63		
	Y79AA1001907	68.02	70.94	96.4	118.34	86.75	104.86		
	Y79AA1001908	0.64	8.4	2.29	1.92	2.59	2.31		
15	Y79AA1001923	1.61	6.64	3.03	3.86	3.76	3.35		
	Y79AA1001927	19.1	22.05	36.94	42.46	45.29	48.81	*	+
	Y79AA1001930	4.07	6.65	8.07	7.92	12.42	12.21		
	Y79AA1001932	2.84	4.41	8.47	11.51	9.1	8.57		
20	Y79AA1001933	2.14	3.27	3.69	4.34	7.99	6.65	*	+
	Y79AA1001942	1.58	3.45	2.69	2.94	2.13	2.41		
	Y79AA1001963	9.6	9.37	46.06	38.48	49.64	47.27		
25	Y79AA1001968	18.61	27.73	37.44	42.93	44.16	55.23	*	+
	Y79AA1001983	1.81	6.35	4.28	3.97	5.86	4.47		
	Y79AA1002000	2.55	5.35	4.55	4.42	3.21	2.83		
	Y79AA1002004	13.1	18.87	27.47	23.72	29.45	40.93		
30	Y79AA1002008	2.51	3.73	3.79	4.54	2.19	2.85		
	Y79AA1002012	1.37	3.22	2.81	3.22	2.29	2.87		
	Y79AA1002017	1.34	2.53	2.46	3.51	3.07	2.82		
35	Y79AA1002022	2.99	4.94	5.93	7.32	7.51	6.01		
	Y79AA1002027	2.02	6.33	2.67	2.69	4.03	4.09		
	Y79AA1002050	2.53	8.12	4.22	6.68	6.91	5.11		
	Y79AA1002058	13.69	21.8	70.12	59.07	70.89	55.33		
40	Y79AA1002060	6.38	13.17	20.54	17.14	21.12	24.23		
	Y79AA1002062	4.33	5.18	8.15	8.54	6.66	5.51		
	Y79AA1002065	33.54	39.97	72.6	49.46	30.04	41.81		
	Y79AA1002067	10.11	11.64	17.24	16.25	9.42	8.13		
45	Y79AA1002069	0.97	1.79	0.54	1.55	1.44	0.66		
	Y79AA1002070	10.16	33.47	44.36	52.16	71.15	73.35	*	+
	Y79AA1002074	38.55	74.38	179.6	165.55	282.48	224.96		
50	Y79AA1002076	0.48	9.71	2.89	2.86	3.34	1.91		
	Y79AA1002083	1.2	7.48	2.03	2.73	1.75	2.06		
	Y79AA1002084	1.79	2.59	4.54	3.73	3.73	2.98		
	Y79AA1002086	0.96	1.78	1.71	2.77	1.88	1.43		
55	Y79AA1002087	11.18	14.9	27.67	33.34	30.01	23.08		

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	Y79AA1001530	7.46	11.69	22	41.43	36.37	36.07	**	+
	Y79AA1001532	5.12	7.35	6.69	10.49	14.82	12.9	*	+
5	Y79AA1001533	1.84	4.89	2.53	4.15	4	4.19		
	Y79AA1001541	2.82	5.89	7.54	7.23	5.08	7.34		
	Y79AA1001548	4.25	4.84	9.82	10.46	7.37	7.1		
10	Y79AA1001555	2	2.55	3.65	4.7	5.28	4.25	*	+
	Y79AA1001562	7.76	10.11	17.15	14.07	16.16	10.83		
	Y79AA1001581	2	5.05	4.47	5.1	7.01	3.54		
	Y79AA1001585	3.18	7.38	10.96	9.72	10.93	8.05		
15	Y79AA1001592	2.61	7.38	5.97	8.15	8	7.02		
	Y79AA1001594	0.76	4.73	3.85	1.96	3.24	1.73		
	Y79AA1001603	56.74	70.81	153.14	131.56	112.16	107.66		
20	Y79AA1001613	3.74	3.52	14.81	13.12	15.36	10.66		
	Y79AA1001630	0.71	2.36	1.73	1.14	2.64	0.67		
	Y79AA1001647	1.96	3.57	2.47	4.14	4.32	2.65		
	Y79AA1001664	4.67	8.39	11.43	8.96	10.01	8.73		
25	Y79AA1001665	1.39	6.4	3.73	4.67	4.71	3.75		
	Y79AA1001679	8.92	15.94	20.71	20.53	26.67	25.35		
	Y79AA1001692	1.87	5.55	3.95	3.99	3.95	3.51		
	Y79AA1001696	1.97	6.49	2.77	1.83	2.63	2.55		
30	Y79AA1001705	6.09	6.44	10.39	7.62	7.92	7.85		
	Y79AA1001711	16.17	12.34	29.74	13.73	23.83	21		
	Y79AA1001717	0.72	2.99	1.29	1.68	3.13	1.14		
35	Y79AA1001719	2.5	5.79	6.44	6.15	6.07	6.43		
	Y79AA1001727	6.87	12.13	14.99	8.73	14.71	8.77		
	Y79AA1001750	10.21	13.63	21.67	21.92	32.29	24.33		
	Y79AA1001760	25.24	27.31	122.97	113.56	155.17	83.24		
40	Y79AA1001777	1.17	3.59	1.6	2.49	1.75	1.28		
	Y79AA1001781	0.31	2.3	0.42	1.62	1.84	1.42		
	Y79AA1001787	1	3.94	3.54	5.51	5.18	5.15		
	Y79AA1001793	16.23	15.19	91.7	60.44	87.21	75.35		
45	Y79AA1001795	1.23	3.84	2.24	2.05	2.45	2.8		
	Y79AA1001799	4.9	8.35	6.99	10.7	10.72	11.26	*	+
	Y79AA1001800	2.25	8.3	10.1	8.49	10.51	9.9		
50	Y79AA1001801	1.77	6.44	4.87	7.67	5.91	7.33		
	Y79AA1001803	0.74	2.15	1.72	1.85	1.68	1.17		
	Y79AA1001805	6.05	6.88	12.15	10.21	8.32	11.15		
	Y79AA1001807	3.37	6.33	12.56	11.76	17.8	16.79		
55	Y79AA1001827	1.7	4.41	3.12	3.43	3.6	2.52		

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	Y79AA1001179	6.81	8.66	16.73	22.82	22.64	20.07	*	+
	Y79AA1001185	1.33	5.3	4.55	3.65	4.49	5.8		
5	Y79AA1001201	5.69	11.3	-16.13	14.57	15.21	19.38		
	Y79AA1001205	1.87	3.28	2.85	5.87	4.85	4.09	*	+
	Y79AA1001211	1.64	4.75	6.93	4.83	4.36	4.15		
10	Y79AA1001212	3.55	6.93	15.91	13.74	15	11.65		
	Y79AA1001216	52.59	51.46	93.73	76.52	97.53	109.55		
	Y79AA1001228	6.1	11.21	9.34	8.99	12.19	10.24		
	Y79AA1001233	0.68	11.46	2.39	0.92	1.66	1.09		
15	Y79AA1001236	4.46	12.86	9.25	11.4	10.66	13.08		
	Y79AA1001239	4.62	13.93	9.94	11.53	12.15	12.94		
	Y79AA1001240	8.74	8.6	13.75	9.13	6.68	3.01		
20	Y79AA1001255	10.37	12.22	22.61	12.47	7.51	6.57		
	Y79AA1001264	3.63	5.15	4.49	7.73	8.59	8.75	**	+
	Y79AA1001272	10.81	13.63	17.47	21.56	20.67	21.32	*	+
	Y79AA1001281	0.45	4.95	1.89	1.42	1.81	0.95		
25	Y79AA1001299	2.49	11.34	9.06	9.9	9.26	9.81		
	Y79AA1001312	2.49	10.36	5.17	2.15	4.77	4.14		
	Y79AA1001319	3.34	11.88	5.27	7.23	6.05	7.15		
	Y79AA1001323	1.22	1.16	2.09	1.11	1.55	0.89		
30	Y79AA1001328	2.04	3.18	3.62	4.66	4.48	4.05	*	+
	Y79AA1001343	154.19	151.55	345.05	304.88	394.54	265.65		
	Y79AA1001351	0.81	2.77	1.67	0.03	1.26	0.96		
35	Y79AA1001364	1.65	6.07	4.03	3.39	4.43	3.6		
	Y79AA1001367	2.16	7.41	2.93	3.09	5.34	3.19		
	Y79AA1001384	0.5	5.14	1.98	0.73	1.15	0.94		
	Y79AA1001391	0.59	3.73	2.88	1.35	1.65	1.2		
40	Y79AA1001394	3.12	4.66	12.92	10.94	9.56	10.94		
	Y79AA1001402	2.77	3.7	5.95	5.65	5.09	4.14		
	Y79AA1001410	0.82	2.78	2.33	2.06	2.31	2.25		
	Y79AA1001414	2.76	7.5	7.59	11.08	10.73	10.06	*	+
45	Y79AA1001426	0.61	4.36	2.61	1.5	1.43	1.82		
	Y79AA1001427	14.22	13.44	86.36	59.92	88.36	63.53		
	Y79AA1001430	11.28	16.98	20.98	29.04	34.03	34.38	**	+
50	Y79AA1001439	16.22	21.53	33.42	45.02	43.74	43.1	*	+
	Y79AA1001485	1.65	2.51	4.38	4.85	3.88	3.57		
	Y79AA1001493	1.29	2.3	3.43	2.43	2.35	2.46		
	Y79AA1001511	4.79	8.57	11.05	9.39	9.11	6.47		
55	Y79AA1001523	2.64	6.57	5.08	8.74	7.37	6.1		

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	Y79AA1000962	1.22	4.45	3.18	2.9	2.41	1.79		
	Y79AA1000963	18.6	26.86	35.93	31.61	42.17	49.13		
5	Y79AA1000966	18.52	19.25	98.18	77.47	116.12	61.1		
	Y79AA1000967	8.62	8.82	33.82	34.47	40.36	29.3		
	Y79AA1000968	3.32	5.67	6.89	8.86	9.4	7.96	*	+
	Y79AA1000969	0.91	3.18	3.04	2.66	2.91	1.41		
10	Y79AA1000976	1.43	4.72	3.02	2.55	3.51	1.43		
	Y79AA1000978	1.99	5.6	7.26	6.9	10.54	5.56		
	Y79AA1000985	9.39	12.84	40.49	38.26	44.3	27.89		
15	Y79AA1000989	21.59	22.49	46.19	51.84	58.65	55.19	*	+
	Y79AA1000991	22.11	22.21	110.42	72.46	96.96	82.23		
	Y79AA1001013	59.2	62.64	140.9	174.85	214.13	201.9	*	+
	Y79AA1001014	2.27	4.16	4.1	4.8	5.2	6.55		
20	Y79AA1001019	3.37	5.89	7.74	9.24	9.02	9.43	*	+
	Y79AA1001020	5.37	7.82	9.43	12.31	11.11	10.86	*	+
	Y79AA1001023	0.83	6.11	2.29	1.22	1.95	1.54		
25	Y79AA1001030	4.23	8.79	10.87	11.14	10.72	12.43		
	Y79AA1001035	0.19	2.88	0.03	14.44	8.19	17.16	*	+
	Y79AA1001041	1.78	2.46	2.36	2.93	2.45	2.78		
	Y79AA1001043	11.65	12.62	15.22	8.64	12.01	14.71		
30	Y79AA1001048	1.1	4.78	3.73	4.05	4.52	4.21		
	Y79AA1001056	4.56	7.82	11.04	8.27	7.11	9.94		
	Y79AA1001061	1.53	7.79	5.28	6.13	7.46	6.66		
35	Y79AA1001062	2.62	6.14	5.02	4.44	6.01	4.67		
	Y79AA1001068	3.46	6.39	7.29	6.61	8.69	7.05		
	Y79AA1001073	8.19	13.08	17.46	24.14	22.1	29.81	*	+
	Y79AA1001077	7.1	7.08	17.15	14.69	14.74	17.08		
40	Y79AA1001078	3.11	8.34	11.07	5.01	12.15	12.92		
	Y79AA1001081	3.59	5.61	4.94	9.62	9.98	10.5	**	+
	Y79AA1001088	27.75	38.61	69.33	93.1	88.97	113.04	*	+
	Y79AA1001089	4.64	7.8	11.92	22.67	22.6	27.73	**	+
45	Y79AA1001090	1.38	4.15	2.2	3.58	2.83	2.35		
	Y79AA1001105	3.7	5.23	15.81	12.52	22.1	13.35		
	Y79AA1001142	8.53	13.38	15.85	14.28	11.42	22.32		
50	Y79AA1001145	2.22	4.68	5.13	4.97	6.26	5.87		
	Y79AA1001162	2.27	2.91	1.62	1.62	4.56	4		
	Y79AA1001167	0.86	2.76	2.38	1.12	2.35	0.77		
	Y79AA1001176	0.57	3.33	1.14	2.02	1.68	0.88		
55	Y79AA1001177	1.21	5.5	2.22	2.35	3.01	1.99		

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	Y79AA1000682	17.99	53.99	93.7	102.53	110.87	118.22		
	Y79AA1000683	1.87	2.66	3.21	4.27	2.59	2.17		
5	Y79AA1000697	21.76	27.52	-43.01	21.93	24.76	27.31		
	Y79AA1000700	5.07	7.1	7.08	7.51	6.93	9.97		
	Y79AA1000702	5.13	14.57	13.31	41.48	56.57	63.15	**	+
	Y79AA1000704	1.34	5.24	1.14	1.5	2.1	1.18		
10	Y79AA1000705	1.86	9.7	6.24	6.06	7.98	6.45		
	Y79AA1000717	6.18	12.39	9.42	9.73	11.1	8.7		
	Y79AA1000722	5.61	8.6	8.65	26.26	34.81	34.31	**	+
15	Y79AA1000724	6.42	9.77	18.55	26.57	21.7	11.95		
	Y79AA1000726	0.77	1.24	0.83	1.46	2.01	0.71		
	Y79AA1000734	2.05	4.46	4.92	3.57	3.3	2.94		
	Y79AA1000748	0.88	4.38	1.77	2.14	2.99	1.56		
20	Y79AA1000750	4.17	8.47	20.58	18.11	20.02	18.62		
	Y79AA1000752	1.25	5.13	2.33	2.23	3.38	3.38		
	Y79AA1000774	2.11	6.23	4.24	4.28	5.79	4.21		
25	Y79AA1000776	1.2	4.37	1.9	2.13	1.83	2.11		
	Y79AA1000777	4.36	5.84	9.63	10.05	6.99	6.01		
	Y79AA1000778	1.72	3.77	3.79	3.87	4.19	1.44		
	Y79AA1000782	2.08	4.18	3.72	3.53	2.89	2.96		
30	Y79AA1000784	7.04	10.01	7.78	13.87	15.58	14.26	**	+
	Y79AA1000794	0.61	5.21	1.88	2.92	1.69	1.23		
	Y79AA1000800	1.59	5.44	3.82	3.38	2.97	3.39		
35	Y79AA1000802	0.64	4.18	1.15	2.52	1.77	2.1		
	Y79AA1000805	2.29	4.03	2.63	2.43	2.11	2.6		
	Y79AA1000814	2.73	3.9	4.14	4.98	7.09	6.76	*	+
	Y79AA1000823	7.91	9.99	12.07	12.02	12.42	6.56		
40	Y79AA1000824	0.98	2.47	1.84	2.75	2.26	0.79		
	Y79AA1000827	1.6	4.02	7.27	6.71	8.91	6.14		
	Y79AA1000831	7.04	10.49	17.32	26.61	30.43	27.82	**	+
	Y79AA1000833	62.14	67.46	191.76	270.42	308.16	248.21	*	+
45	Y79AA1000850	1.69	5.68	2.72	4.92	4.59	4.21		
	Y79AA1000856	3.49	6.78	6.31	9.28	6.01	8.51		
	Y79AA1000862	2.22	2.76	2.44	3.08	2.87	2.59		
50	Y79AA1000876	7.46	10.04	17.91	27.36	27.61	25.46	**	+
	Y79AA1000888	4.59	5.07	28.1	24.51	38.78	22.48		
	Y79AA1000902	4.65	5.74	8.44	12.18	12.32	7.68		
	Y79AA1000935	3.53	5.99	6.69	8.28	10.07	9.18	*	+
55	Y79AA1000959	0.74	6.29	4.35	6.71	5.77	6.07		

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	Y79AA1000342	10.65	10.05	26.47	23.7	19.13	28.35		
	Y79AA1000346	7.61	8.17	7.9	20.1	22.06	20.18	**	+
5	Y79AA1000347	6.94	7.96	12.42	18.78	16.47	18.48	**	+
	Y79AA1000349	6.93	9.63	12.67	12.31	11.96	14.75		
	Y79AA1000355	3.17	8.28	8.94	8.84	13.55	8.77		
	Y79AA1000368	5.24	8.39	24.43	22.48	35.67	22.55		
10	Y79AA1000388	22.9	32.66	62.37	94.17	128.35	109.08	**	+
	Y79AA1000392	3.02	6.81	3.76	3.42	2.73	3.78		
	Y79AA1000405	3.98	5.97	8.25	7.14	8.17	9.32		
15	Y79AA1000410	6.01	7.87	15.72	13.79	17.05	14.95		
	Y79AA1000420	1.54	4.78	3.13	3.32	3.95	5.1		
	Y79AA1000423	1.38	7.08	5.59	5.22	6.04	10.27		
	Y79AA1000426	3.61	9.44	8.66	4.24	4.43	5.22		
20	Y79AA1000432	0.8	4.79	2.16	1.91	2.01	2.34		
	Y79AA1000453	23.94	30.67	47.79	39.74	50.65	58.24		
	Y79AA1000465	4.12	6.02	6.65	4.77	4.14	7.69		
25	Y79AA1000469	11.59	9.61	18.04	13.82	16.21	17.18		
	Y79AA1000480	1.24	4.37	2.78	3.33	3.57	2.79		
	Y79AA1000502	5.31	7.97	12.58	10.49	11.35	15.26		
	Y79AA1000521	1.24	4.4	4.13	2.51	3.61	2.7		
30	Y79AA1000534	3.22	8.13	8.92	11.97	14.41	13.46	*	+
	Y79AA1000538	3.58	6.95	8.79	9.52	12.12	8.41		
	Y79AA1000539	12.76	14.96	53.11	42.61	68.56	50.97		
	Y79AA1000540	1.32	3.59	1.61	2.54	2.97	3.21		
35	Y79AA1000560	160.46	140.99	339.33	380.8	313.21	220.43		
	Y79AA1000574	1	2.92	1.65	1.98	2.04	1.59		
	Y79AA1000584	2.07	4.55	4.97	4.62	5.39	4.04		
40	Y79AA1000589	10.74	13.67	81.43	59.09	95.35	68.5		
	Y79AA1000598	1.43	7.64	2.17	1.85	3.88	3.18		
	Y79AA1000600	2.7	10.02	7.93	13.64	15.64	12.84	*	+
	Y79AA1000609	1.18	5.16	1.44	2.28	2.77	1.55		
45	Y79AA1000618	1.85	10.59	5.76	7.4	9.5	9.37		
	Y79AA1000627	1.91	3.93	4.57	3.27	3.02	2.43		
	Y79AA1000636	5.16	5.7	9.9	15.57	11.52	5.38		
50	Y79AA1000649	9.45	10.97	12.73	18.7	11.56	20.54		
	Y79AA1000656	15.32	20.21	96.75	80.17	115.97	82.61		
	Y79AA1000673	1.02	5.86	2.14	1.39	3.13	2.4		
	Y79AA1000674	11.88	21.96	78.28	59.4	98.22	62.67		
55	Y79AA1000678	2.48	8.91	3.88	3.01	4.15	2.45		

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	Y79AA1000030	1.79	4.06	4.52	3.24	3.85	2.47		
	Y79AA1000033	2.87	6	8.4	8.37	10.17	5.83		
5	Y79AA1000037	1.38	3.36	-5.71	4.84	6.82	4.49		
	Y79AA1000041	1.05	5.16	3.79	4.73	3.65	2.06		
	Y79AA1000059	1.69	5	4.09	3.88	4.51	2.82		
10	Y79AA1000065	24.06	28.99	52.25	82.48	101.48	98.73	**	+
	Y79AA1000081	39.47	49.78	73.62	113.19	114.49	98.22	**	+
	Y79AA1000127	4.08	4.21	5.8	8.42	10.03	7.39	*	+
	Y79AA1000130	2.24	2.48	5.76	6.61	8.1	8.03	*	+
15	Y79AA1000131	507.64	569.21	946.04	769.75	725.35	342.07		
	Y79AA1000134	1.99	4.93	4.21	5.63	4.75	5.38		
	Y79AA1000143	3.58	8.79	4.83	9.98	10.98	11.04	*	+
	Y79AA1000144	4.63	10.79	10.59	11.02	11.62	11		
20	Y79AA1000150	18.39	22.18	84.69	93.5	117.62	78.89		
	Y79AA1000153	183.67	191.4	436.64	423.46	442.52	386.45		
	Y79AA1000166	2.13	2.25	4.15	3.52	4.97	3.6		
25	Y79AA1000179	2.58	3.76	4.2	6.85	7.58	3.89		
	Y79AA1000181	1.96	3.92	4.2	4.82	5.66	3.79		
	Y79AA1000202	22.93	24.47	55.57	91.68	86.86	83.22	**	+
	Y79AA1000207	5.22	7.51	9.82	14.95	16.24	12.62	*	+
30	Y79AA1000214	14.94	22.18	33.76	50.43	60.78	41.96	*	+
	Y79AA1000222	11.8	14.89	21.69	49.21	58.01	68.86	**	+
	Y79AA1000226	11.04	14.94	34.41	22.86	33.54	30.03		
35	Y79AA1000227	5.95	4.52	7.25	8.98	9.51	9.62	*	+
	Y79AA1000230	1.09	1.49	2.02	2.07	2.88	2.59		
	Y79AA1000231	5.99	9.04	15.81	14.63	23.77	17.1		
	Y79AA1000239	15.47	20.55	25.65	18.95	24.01	22.11		
40	Y79AA1000258	2.64	5.17	6.72	5.87	4.95	5.84		
	Y79AA1000268	2.65	5.48	5.09	4.33	5.76	4.01		
	Y79AA1000269	4.32	7.88	7.86	8	6.86	8.24		
45	Y79AA1000270	5.28	8.35	11.58	13.17	17.58	16.23	*	+
	Y79AA1000280	1.74	4.17	4.9	5.29	3.1	5.27		
	Y79AA1000285	3.44	4.21	5.91	4.01	6.86	5.22		
	Y79AA1000295	0.75	3.06	4.85	4.32	4.23	4.45		
50	Y79AA1000307	2.88	3.91	5.06	9.35	7.58	11.25	*	+
	Y79AA1000313	3.11	9.02	9.85	10.61	12.84	13.36		
	Y79AA1000314	4.23	10.74	9.19	6.93	6.53	7.51		
	Y79AA1000328	4.65	10.05	2.64	7.73	9.28	8.68		
55	Y79AA1000334	1.43	4.22	3.55	2.68	2.81	3.46		

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	VESEN1000669	8.43	10.02	16.35	15.06	14.17	15.51		
	VESEN1000743	0.97	3.3	2.52	1.99	3.37	3.21		
5	VESEN1000752	37.43	51.51	-72.35	49.32	57.03	57.96		
	VESEN1000761	13.48	18.17	19.37	24.6	33.21	31.28	*	+
	VESEN2000039	10.45	15.98	15.56	13.56	18.3	20.1		
10	VESEN2000102	0.4	3.99	1.6	1.21	1.51	1.61		
	VESEN2000164	2.45	3.52	4.8	5.55	4.36	3.66		
	VESEN2000175	0.57	2.64	1.94	3.03	2.05	2.59		
	VESEN2000186	3.77	5.53	6.53	6.68	3.87	2.47		
15	VESEN2000199	8.94	13.26	21.75	19.58	24.45	24.12		
	VESEN2000200	0.5	4.97	2.78	3.03	3.1	1.6		
	VESEN2000204	0.48	12.7	1.02	0.98	1.2	0.33		
20	VESEN2000218	6.66	20.26	19.48	21.37	20.75	18.86		
	VESEN2000230	0.84	7.4	1.45	2.74	1.7	2.41		
	VESEN2000272	2.29	4	8.92	7	8.31	5.88		
	VESEN2000299	1.99	2.97	3.2	3.3	3.77	3.31		
25	VESEN2000323	4.51	8.12	8.37	8.91	9.5	9.36		
	VESEN2000327	3.16	5.8	4.42	3.62	7.66	5.07		
	VESEN2000328	5.44	8.02	11.88	8.73	15.35	14.36		
	VESEN2000330	6.39	15.42	14.7	14.59	27.73	18.86		
30	VESEN2000336	0.82	8.97	2.54	3.46	3.83	2.88		
	VESEN2000354	1.56	8.24	2.48	1.53	2.71	2.1		
	VESEN2000378	7.17	8.87	14.57	13.23	11.14	10.3		
35	VESEN2000379	19.87	23.02	44.55	49.13	42.81	32.61		
	VESEN2000397	0.72	2.38	1.24	1.36	2.06	1.54		
	VESEN2000416	2.83	3.88	4.41	5.74	5.31	5.71	*	+
	VESEN2000420	1.08	3.58	1.94	0.95	1.21	1.13		
40	VESEN2000430	0.51	6.68	1.53	2.06	1.79	1.8		
	VESEN2000448	0.51	6.87	1.73	2.12	2.69	1.5		
	VESEN2000449	2.43	8.07	6.59	8.3	11.31	8.21		
45	VESEN2000456	0.74	0.87	2.11	1.54	1.22	0.87		
	VESEN2000562	4.07	3.42	17.42	13.67	22.82	14.47		
	VESEN2000573	0.18	1.75	1.75	1.04	1.21	1.13		
	VESEN2000604	1.73	3.44	2.24	2.04	1.67	2.15		
50	VESEN2000614	4.16	9.02	14.64	13.54	16.27	12.27		
	VESEN2000638	0.48	5.92	1.98	1.33	1.91	1.68		
	VESEN2000641	0.83	3.69	1.34	1.95	2.21	1.43		
	VESEN2000645	2.18	5.29	5.38	5.91	5.73	5.5		
55	Y79AA1000013	2.57	2.7	4.33	3.45	3.8	3.94		

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	THYRO1001907	2.16	3.08	3.37	3.9	4.53	2.74		
	TRACH1000006	2.51	5.87	6.63	4.34	3.46	3.22		
5	TRACH1000013	1.53	4.65	3.68	2.55	3.33	2.65		
	TRACH1000074	2.65	6.75	6.09	7.5	7.26	4.77		
	TRACH1000095	0.28	5.66	2.23	2.46	1.48	1.11		
	TRACH1000102	2.42	6.66	4.09	5.79	5.04	3.65		
10	TRACH1000108	1.1	1.01	2.05	1.75	2.49	1.09		
	TRACH1000126	0.96	1.75	2.71	1.82	3.79	2.54		
	TRACH1000146	1.3	2.67	2.31	3.02	5.18	3.75		
15	TRACH1000160	0.61	4.06	1.5	1.47	1.76	0.72		
	TRACH1000184	4.45	7.16	10.16	7.47	8.73	5.69		
	VESEN1000004	0.69	5.55	3.19	2.56	2.95	2.02		
	VESEN1000007	0.93	5.32	2.94	2.38	3.45	2.94		
20	VESEN1000013	5.96	10.11	16.78	10.76	11.25	13.88		
	VESEN1000028	5.2	7.5	9.88	13.18	11.71	14.08	*	+
	VESEN1000059	1.55	2.88	2.1	3.38	2.82	2.27		
25	VESEN1000100	1.96	3.22	3.35	3.49	4.58	3.59		
	VESEN1000107	0.88	4.84	2.88	3.12	2.9	2.48		
	VESEN1000117	1.63	6.43	2.46	2.16	2.7	1.79		
	VESEN1000122	1.52	5.34	1.24	4.79	4.51	4.5		
30	VESEN1000137	0.76	5.47	1.92	1.75	3.33	1.65		
	VESEN1000195	7.79	7.93	11.67	8.42	7.51	10.27		
	VESEN1000215	1.48	3.03	2.06	2.67	3.84	1.87		
35	VESEN1000279	8.71	11.32	18.49	22.93	23.38	34.68	*	+
	VESEN1000363	3.52	6.07	9.99	7.2	9.06	4.59		
	VESEN1000388	2.55	6.48	3.31	4.17	3.75	6.7		
	VESEN1000394	0.44	7.11	2.33	2.37	2.55	2.36		
40	VESEN1000410	1.11	5	1.78	2.36	2.71	3.69		
	VESEN1000411	2.37	4.95	5.08	6.76	7.55	9	*	+
	VESEN1000415	1.54	2.64	4.03	5.57	3.92	5.29		
45	VESEN1000440	7	5.53	7.81	3.79	9.4	12.22		
	VESEN1000452	1.22	3.65	2.33	2.91	3.97	4.11		
	VESEN1000539	191.54	185.28	334.6	389.84	403.89	547.31	*	+
	VESEN1000554	0.67	6.47	1.43	1.47	2.55	1.58		
50	VESEN1000557	4.22	7.94	7.73	6.55	9.07	10		
	VESEN1000575	7.49	9.75	16.33	11.95	11.73	14.8		
	VESEN1000585	1.69	4.49	3.37	2.53	2.93	3.08		
	VESEN1000592	1.58	2.31	1.58	2.02	1.83	1.46		
55	VESEN1000658	1.96	3.56	4.45	5.86	3.91	4.91		

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	THYR01001602	1.49	3.26	3.52	4.95	5.22	3.41		
	THYR01001605	1.58	4.48	3.22	3.2	3.43	2.42		
5	THYR01001608	1.87	9.45	5.1	5.04	8.23	4.7		
	THYR01001617	6.06	13.68	11.47	9.75	10.87	9.61		
	THYR01001634	1.87	9.08	3.46	2.93	5.59	2.05		
	THYR01001637	3.51	3.13	9.65	8.72	7.94	9.07		
10	THYR01001641	2.57	3.73	5.09	4.03	3.08	2.94		
	THYR01001656	1.59	2.94	4.16	2.82	5.36	2.33		
	THYR01001658	22.34	29.19	40.11	34.98	33.16	42.01		
15	THYR01001661	1.4	5.83	2.31	2.93	3.31	2.05		
	THYR01001671	0.67	7.36	2.68	1.89	1.34	1.8		
	THYR01001672	1.1	9.24	2.1	1.14	1.52	1.66		
	THYR01001673	1.59	7.6	3.49	2.86	4.74	2.16		
20	THYR01001677	1.6	2.27	3.87	3.03	3.54	3.36		
	THYR01001683	12.71	17.66	29.06	24.4	15.4	16.72		
	THYR01001700	1.39	2.52	2.67	2.09	1.58	1.37		
25	THYR01001702	11.83	15.98	16.19	15.63	14.35	14.29		
	THYR01001703	1.63	6.74	4.25	4.72	3.27	4.21		
	THYR01001706	1.7	6.47	3.01	2.96	5.6	3.53		
	THYR01001721	1.84	5.66	3.2	2.73	6.37	2.77		
30	THYR01001725	5.3	6.55	9.69	8.97	8.65	8.29		
	THYR01001730	17.72	20.4	40.1	30.61	26.56	34.8		
	THYR01001738	1.35	3.18	4.65	3.52	2.82	1.78		
	THYR01001743	0.19	2.13	1.85	1.8	1.64	1.06		
35	THYR01001745	0.47	2.88	1.55	1.05	1.2	1.27		
	THYR01001746	1.9	6.25	4.04	6.12	4.01	3.88		
	THYR01001770	15.49	20.38	35.39	41.65	44.42	40.17	*	+
40	THYR01001772	1.12	4.88	3.64	4.78	4.24	3.06		
	THYR01001778	3.89	6.68	9.89	14.67	13.47	14.25	*	+
	THYR01001793	3.85	3.77	9.43	10.3	10.42	4.92		
	THYR01001796	1.35	2.28	2.28	3.45	4.22	3.24	*	+
45	THYR01001800	1.82	2.99	2.75	4.17	5.12	2.09		
	THYR01001803	3.42	6.03	5.21	4.31	4.14	3.42		
	THYR01001809	1.6	4.26	3.4	5.9	4.23	3.39		
50	THYR01001817	8.69	18.33	24.88	22.11	22.77	25.27		
	THYR01001819	4.68	8.46	9.01	7.84	10.46	6.77		
	THYR01001828	21.89	24.53	104.32	106.68	121.95	68.55		
	THYR01001854	6.67	6.12	12.45	13.43	15.86	13.31		
55	THYR01001895	0.85	1.31	2.52	3.54	4.31	1.28		

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	THYRO1001401	1.76	5.33	4.89	5.39	7.86	6.29		
	THYRO1001403	1.26	5.15	3.22	3.22	4.42	3.94		
5	THYRO1001405	6.99	12.5	-10.86	6.69	8.56	10.63		
	THYRO1001406	15.73	14.87	27.69	21.36	22.77	21.36		
	THYRO1001411	4.49	5.46	10.08	8.93	12.44	8.12		
	THYRO1001420	11.55	15.25	47.52	42.01	44.49	49.87		
10	THYRO1001426	3.42	5.56	8.83	9.32	12.77	11.18		
	THYRO1001430	6.97	6.54	10.84	11.13	11.7	13.81		
	THYRO1001434	0.68	5.19	2.11	2.08	4.23	1.73		
15	THYRO1001456	1.74	6.05	2.63	2.89	2.66	2.4		
	THYRO1001457	1.71	4.72	2.04	2.95	4.7	2.67		
	THYRO1001458	0.95	5.44	6.11	6.13	9.17	7.23		
	THYRO1001459	4.54	5.07	9.42	7.18	9.87	14.21		
20	THYRO1001471	0.91	2.07	1.93	2.36	2.91	1.64		
	THYRO1001478	0.58	3.09	1.34	0.95	2.61	2.75		
	THYRO1001480	5.4	10.53	13.62	14.79	15.94	15.57		
25	THYRO1001481	2.95	8.64	7.24	4.91	7.76	7.13		
	THYRO1001487	1.36	5.51	3.52	4.1	2.67	3.48		
	THYRO1001495	2.06	5.57	5.43	5.06	9.4	7.55		
	THYRO1001498	5.39	8.08	13.42	11.59	16.38	19.6		
30	THYRO1001510	1.67	2.88	1.59	1.65	2.57	3.35		
	THYRO1001512	26.7	26.97	110.28	56.03	90.95	68.54		
	THYRO1001519	5.92	7.77	11.05	8.91	5.75	10.38		
35	THYRO1001522	2.02	4.7	5.79	4.15	4.69	5.51		
	THYRO1001523	1.83	4.92	3.73	4.38	4.07	4.26		
	THYRO1001526	26.21	28.22	44.73	34.28	47.64	46.52		
	THYRO1001529	1.64	5.27	2.8	2.32	2.42	2.98		
40	THYRO1001534	1.41	4.01	4.96	5.1	5.9	4.54		
	THYRO1001537	7.4	5.17	12.33	5.97	7.42	7.23		
	THYRO1001541	2.14	3.88	8.27	7.76	8.7	6.73		
	THYRO1001545	1.26	3.84	2.9	4.95	3.57	3.16		
45	THYRO1001559	4.52	6.34	8.04	9.06	10.5	10.54	*	+
	THYRO1001563	9.49	14.06	15.89	10	15.49	22.09		
	THYRO1001570	2.01	8.2	3.85	4.25	5.17	3.41		
50	THYRO1001573	1.15	5.77	2.22	1.47	2.87	2.67		
	THYRO1001584	2.47	8.54	8.38	5.14	7.81	7.29		
	THYRO1001593	4.27	5.67	11.17	9.5	10.93	9.52		
	THYRO1001595	3.14	4.53	7.06	5.97	6.35	7.29		
55	THYRO1001596	4.71	5.48	7.44	6.45	5.86	2.51		

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	THYR01001109	1.81	6.02	2.68	3.06	2.58	1.99		
	THYR01001113	11.41	17.42	32	21.81	26.65	18.72		
5	THYR01001120	1.65	6.22	5.27	4.78	5.8	3.72		
	THYR01001121	1.57	4.28	4.19	3.92	2.72	3		
	THYR01001128	1.64	2.77	5.86	3.52	3.19	5.09		
	THYR01001133	1.14	3.02	7.23	6.54	4.54	4.12		
10	THYR01001134	2.97	4.78	1.63	3.14	2.83	1.38		
	THYR01001142	0.3	2.69	0.63	1.3	1.71	0.22		
	THYR01001173	8.37	12.87	7.72	11.14	9.92	10.62		
15	THYR01001175	3.26	6.63	5.51	3.46	4.62	3.52		
	THYR01001177	1.36	5.85	5.93	4.66	7.27	7.97		
	THYR01001189	2.74	6.93	11.42	7.84	7.27	9.94		
	THYR01001194	1.05	2.62	4.96	4.89	4.57	2.31		
20	THYR01001204	2.17	3.58	4.27	4.03	4.74	4		
	THYR01001205	5.76	10.65	20.23	18.54	19.57	20.9		
	THYR01001213	1.21	4.69	4.44	3.12	3.21	2.33		
25	THYR01001224	3.59	8.25	6.37	9.92	12.69	10.55	*	+
	THYR01001237	2.82	6.25	4.99	3.61	4.53	4.46		
	THYR01001242	9.74	11.65	19.04	20.02	19.98	20.46		
	THYR01001258	2.08	5.45	3.58	3.33	2.05	2.66		
30	THYR01001262	0.86	2.64	3.38	2.36	3.61	2.69		
	THYR01001266	0.15	2.39	1.02	0.97	1.64	0.66		
	THYR01001271	1.85	4.12	4.12	2.46	2.77	2.97		
	THYR01001287	7.3	8.3	39.26	30.14	43.68	26.2		
35	THYR01001290	0.38	3.25	1.14	1.15	1.35	0.36		
	THYR01001291	0.96	7.17	4.38	4.31	4.97	3.5		
	THYR01001297	3.05	8.04	6.14	6.85	7.47	9.18		
40	THYR01001302	1.72	5.59	5.17	3.8	3.71	3.5		
	THYR01001313	1.61	2.33	2.91	2.91	2.62	1.48		
	THYR01001320	1.76	2.52	5.31	5.07	5.74	4.83		
	THYR01001321	2.25	2.65	4.3	2.48	4.23	4.23		
45	THYR01001322	1.34	3.93	3.34	1.75	2.67	2.01		
	THYR01001327	1.29	6.01	4.18	1.4	3.89	2.49		
	THYR01001336	1.89	6.84	6.72	4.62	4.43	4.18		
	THYR01001347	0.43	4.12	3.35	1.85	2.81	0.65		
50	THYR01001358	2.57	5.74	4.52	4.3	5.75	5.1		
	THYR01001363	0.8	2.15	1.52	2.09	2.24	2.28		
	THYR01001365	0.86	3	1.6	2.19	2.6	1.96		
55	THYR01001374	1.85	4.45	12.86	9.4	13.01	6.21		

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	THYRO1000852	1.59	6.02	5.63	2.8	4.7	3.32		
	THYRO1000855	3.14	5.02	6.63	9.03	15.1	10.07	*	+
5	THYRO1000865	1.86	4.3	11.97	10.01	11.47	8.95		
	THYRO1000866	7.47	6.29	12.66	4.49	7.87	6.01		
	THYRO1000881	5.62	7.3	10.93	15.65	26.64	29.58	*	+
10	THYRO1000894	0.33	3.95	1.36	1.75	1.48	1		
	THYRO1000895	0.58	4.43	1.42	1.62	1.46	0.82		
	THYRO1000916	1.22	5.49	3.43	2.43	3.13	2.29		
	THYRO1000917	16.19	25.26	34.11	30.37	37.42	35.89		
15	THYRO1000926	0.78	3.13	1.27	1.76	1.57	0.82		
	THYRO1000934	0.08	3.1	1.34	0.43	1.38	1.46		
	THYRO1000951	0.52	2.46	1.26	2.33	2.11	1.9		
20	THYRO1000952	2.25	3.81	6.01	2.38	2.53	2.24		
	THYRO1000956	0.06	2.55	1.81	1.16	1.5	0.87		
	THYRO1000960	0.5	6.72	2.89	1.85	2.79	1.48		
	THYRO1000961	1.67	7.77	3.56	4.73	5.26	4.64		
25	THYRO1000964	0.42	11.59	0.76	1	1.27	1.06		
	THYRO1000971	1.82	9.9	3.56	3.29	3.33	2.19		
	THYRO1000974	2.87	8.83	7.53	9.87	11.79	8.71		
	THYRO1000975	1.5	2.19	3.8	4.02	3.68	3.68		
30	THYRO1000983	6.42	8.31	11.63	12.67	8.49	7.12		
	THYRO1000984	2.4	2.83	3.03	3.29	2.98	3.26		
	THYRO1000988	1.36	4.14	3.23	3.48	3.68	2.67		
35	THYRO1000991	1.22	4.71	2.05	1.76	2.22	3.2		
	THYRO1000999	0.87	9.64	3.26	1.96	3.14	2.26		
	THYRO1001003	2.97	8.43	4.42	3.1	4.52	3.56		
	THYRO1001015	0.6	6.29	2.04	2.22	1.79	1.66		
40	THYRO1001016	1.73	2.26	3.34	2.06	1.85	1.24		
	THYRO1001022	0.9	1.86	0.86	1.68	1.25	1.41		
	THYRO1001031	4.65	3.97	4.55	5.03	7.03	6.16		
45	THYRO1001033	1.18	3.34	2.46	2.86	3.45	2.02		
	THYRO1001062	1.21	5.4	4.14	2.9	4.31	2.62		
	THYRO1001063	0.5	8.74	2.38	2.37	2.09	2.84		
	THYRO1001071	0.12	7.45	0.88	1.33	0.68	0.76		
50	THYRO1001080	2.56	6.75	5.11	4.96	4.31	4.78		
	THYRO1001093	0.77	1.63	3.24	5.11	1.74	1.5		
	THYRO1001100	0.52	1.89	2.05	1.89	1.21	0.78		
	THYRO1001102	2.61	3.6	5.7	4.4	4.95	6.93		
55	THYRO1001104	3.67	6.54	6.55	8.77	8.01	11.18		

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	THYR01000637	0.91	3.96	1.71	1.18	2.03	1.54
	THYR01000641	0.38	4.19	2.49	1.36	1.67	1.64
5	THYR01000657	2.99	3.69	5.42	7.67	12.28	3.86
	THYR01000658	2.68	3.62	5.39	5.4	5.55	6.09
	THYR01000662	1.1	3.19	2.09	2.42	2.69	1.66
10	THYR01000666	0.57	3.19	2.28	1.63	1.48	1.43
	THYR01000676	1.37	4.53	2.01	1.75	1.83	1.56
	THYR01000678	0.52	5.86	0.99	1.29	1.4	0.53
	THYR01000684	0.95	4.98	2.94	1.92	2.65	1.47
15	THYR01000694	2.08	6.64	4.65	2.8	2.48	3.59
	THYR01000699	2.98	2.14	5.55	4.86	7.08	7.12
	THYR01000712	1.88	4.25	5.9	6.25	6.75	7.78
20	THYR01000715	5.74	5.67	27.37	21.74	28.63	16.99
	THYR01000716	0.92	3.26	3.2	1.88	1.78	1.35
	THYR01000717	1.58	5	4.36	2.98	4.63	1.91
	THYR01000723	0.6	4.54	1.6	0.55	1.06	0.85
25	THYR01000734	-0.01	4.81	1.89	1.49	1.73	1.07
	THYR01000748	0.98	5.51	5.23	2.35	3.85	3.18
	THYR01000755	1.74	3.26	4.32	4.33	3.47	4.38
	THYR01000756	2.79	4.24	3.24	3.46	4.2	3.41
30	THYR01000776	0.48	2.17	3.02	3.36	3.99	3.34
	THYR01000777	1.81	3.39	4.54	4.99	2.05	2.37
	THYR01000779	1.45	3.55	0.88	0.18	1.01	-0.26
35	THYR01000782	3.92	10.13	12.52	10.76	15.05	14.05
	THYR01000783	0.12	5.51	1.2	1.11	1.41	0.92
	THYR01000786	6.65	9.54	19.71	15.74	7.92	13.7
	THYR01000787	0.23	1.88	1.67	1.31	1.54	0.78
40	THYR01000792	1.51	3.13	2.29	3.09	3.13	2.11
	THYR01000793	0.11	3.13	0.84	1.51	1.86	1.16
	THYR01000795	1.23	6.03	3.54	2.76	3.1	3.05
	THYR01000796	0.6	7.73	2.44	2.26	2.95	1.66
45	THYR01000798	1.89	5.82	2.51	2.59	3.57	3.53
	THYR01000800	9.26	17.2	24.74	17.74	20.68	21.06
	THYR01000805	0.49	3.04	1.08	0.72	2.66	1.38
50	THYR01000815	2.54	3.49	9.48	7.61	5.47	7.87
	THYR01000829	5.55	7.83	10.57	3.78	8.32	10.01
	THYR01000835	0.96	3.2	1.93	1.07	2.36	1.8
	THYR01000843	1.09	11.48	3.56	3.69	4.41	3.62
55	THYR01000846	0.76	5.71	1.32	2.67	1.62	1.26

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	THYRO1000345	1.36	7.34	11.92	7.82	5.84	9.49		
	THYRO1000358	1.82	3.39	3.08	1.92	2.32	1.54		
5	THYRO1000368	0.76	2.39	-2.73	1.43	2.82	0.58		
	THYRO1000375	3.2	7.03	4.79	7.38	6.09	9.77		
	THYRO1000381	0.92	2.88	2.19	3.87	3.11	2.74		
	THYRO1000387	0.98	6.66	3.22	2.53	3.56	2.51		
10	THYRO1000394	1.31	9.88	4.59	4.29	5.19	2.61		
	THYRO1000395	0.8	10.44	2.26	1.97	2.07	2.02		
	THYRO1000400	0.57	8.1	2.82	2.35	2.96	2.52		
15	THYRO1000401	0.86	1.94	2.5	1.87	1.16	1.57		
	THYRO1000407	1.97	2.3	1.36	1.37	1.58	0.55		
	THYRO1000420	1.8	2.67	4.46	3.52	3.53	3.39		
	THYRO1000438	1.78	4.37	3.26	2.94	3.33	3.15		
20	THYRO1000452	2.62	7.99	6.45	3.71	5.75	4.38		
	THYRO1000455	0.32	6.67	2.31	0.25	0.97	0.87		
	THYRO1000471	0.99	8.03	2.05	1.11	2.08	1.02		
25	THYRO1000481	1.33	6.23	4.68	3.79	3.45	4.55		
	THYRO1000484	1.2	1.42	2.41	2.35	3	2.21		
	THYRO1000488	1.18	2.64	2.44	1.49	2.02	1.7		
	THYRO1000501	1.12	4.01	2.78	3	1.92	1.82		
30	THYRO1000502	0.34	3.7	1.69	1.79	1.44	1.2		
	THYRO1000505	0.13	4.64	1.19	1.14	1.02	0.6		
	THYRO1000535	11.1	20.54	39.24	54.13	69.59	62.96	*	+
35	THYRO1000556	1.89	6.36	4.13	3.77	5.17	3.69		
	THYRO1000558	0.25	2.82	1.12	1.16	0.81	0.61		
	THYRO1000569	2.88	4.12	6.05	5.78	4.46	4.88		
	THYRO1000570	2.31	3.28	8.46	8.53	6.04	3.49		
40	THYRO1000572	0.43	2.04	1.11	0.17	0.97	-0.42		
	THYRO1000573	0.69	4.02	1.73	2.02	2.2	1.78		
	THYRO1000577	1.06	5	1.34	0.96	1.22	0.71		
	THYRO1000580	0.79	3.72	3.01	2.82	2.2	1.79		
45	THYRO1000584	2.18	6.88	8.8	7.57	6.61	7.58		
	THYRO1000585	4.83	9.37	9.83	5.76	6.27	9.52		
	THYRO1000596	0.22	0.93	1.19	0.44	1.36	0.21		
50	THYRO1000602	2.08	2.95	4.01	4.05	4.65	4.97		
	THYRO1000605	0.37	3.01	0.98	2.13	2.14	1.56		
	THYRO1000615	1.02	3.62	1.24	1.55	1.36	1.29		
	THYRO1000625	0.71	5.48	2.28	2.46	2.9	1.78		
55	THYRO1000636	3.67	5.65	6.9	6.53	7.84	6.67		

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	THYR01000092	2.32	5.1	4.66	3.75	4.43	4.5		
	THYR01000093	0.35	3.24	0.83	1.54	1.27	0.95		
5	THYR01000099	0.45	2.53	2.73	2.8	1.67	2.39		
	THYR01000107	0.5	2.95	2.7	2.86	3.22	2		
	THYR01000111	0.85	4.58	1.78	1.4	2.06	2.36		
	THYR01000121	1.33	5.72	2.52	1.94	2.4	2.95		
10	THYR01000124	0.27	5.55	0.64	0.86	0.89	0.64		
	THYR01000129	0.36	2.1	0.11	0.94	1.11	0.92		
	THYR01000130	1.82	3.11	3.13	3.85	3.01	2.39		
15	THYR01000132	2.4	3.62	9.43	11.14	6.99	6.26		
	THYR01000134	1.5	4.07	3.22	4.06	3.65	3.73		
	THYR01000144	1.72	4.78	3.15	7.87	7.09	2.33		
	THYR01000155	1.6	4.1	1.45	1.77	1.9	2.23		
20	THYR01000156	1.13	6.53	3.62	2.45	4.29	2.58		
	THYR01000163	3.62	8.42	5.28	4.76	6.63	2.24		
	THYR01000173	1.19	4.45	2.26	3.33	1.36	2.75		
25	THYR01000186	1.98	3.24	7.86	6.91	6.84	6.35		
	THYR01000187	2.7	3.58	5.3	4.92	6.24	5.22		
	THYR01000190	1.12	3.32	2.94	3.73	4.55	2.71		
	THYR01000196	0.3	5.28	0.81	0.66	1.21	0.52		
30	THYR01000197	2.05	7.28	4.69	4.08	6.24	3.89		
	THYR01000199	0.76	6.28	4.13	1.93	2.08	1.98		
	THYR01000206	8.47	6.92	9.25	8.44	11.6	7.5		
35	THYR01000221	1.9	3.17	4.42	4.02	5.87	4.54		
	THYR01000222	3.65	4.26	4.23	4.68	4.96	4.93	*	+
	THYR01000228	0.81	3.67	2.85	2.24	3.04	2.94		
	THYR01000241	1.76	3.7	6.29	4.62	5.54	4.01		
40	THYR01000242	0.63	4.16	4.46	2.49	2.56	2.62		
	THYR01000246	1.61	5.5	3.9	3.43	4.7	3.91		
	THYR01000253	1.07	4.05	1.73	1.99	3.35	2.31		
45	THYR01000270	1.15	5.12	1.39	1.22	2.5	1.26		
	THYR01000279	0.42	2.84	0.25	0.65	1.01	0.58		
	THYR01000285	2.75	4.65	7.31	7.03	7.75	4.88		
	THYR01000288	7.76	7.59	11.77	5.68	5.07	7.22		
50	THYR01000296	4.18	6.04	6.22	7.4	11.24	8.96	*	+
	THYR01000320	1.54	5.83	4.97	3.65	4.45	3.34		
	THYR01000322	1.1	5.48	2.48	1.76	3.93	1.76		
	THYR01000327	1.75	7.69	4.77	6.21	5.23	4.41		
55	THYR01000343	2.5	6.12	5.35	5.06	5.04	6.13		

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		SPLEN1000166	0.4	2.9	2.96	1.67	1.19	1.68		
		SPLEN1000175	2.16	4.48	6.1	5.65	4.12	4.15		
5		SPLEN1000182	0.98	2.66	-0.23	0.83	0.6	0.67		
		SPLEN1000185	3.41	8.49	8.54	11.38	10.43	11.95		
		THYMU1000004	10.22	14.07	20.43	22.34	22.76	23.6		
		THYMU1000009	9.48	10.13	14.9	13.48	23.86	22.1		
10		THYMU1000015	8.87	10.42	16.18	19.25	22.21	20.8	*	+
		THYMU1000016	6.24	5.96	13.03	10.3	8.45	9.38		
		THYMU1000023	0.77	1.86	3.6	5.22	3.68	3.6		
15		THYMU1000034	0.16	1.77	1.8	0.79	0.88	0.14		
		THYMU1000035	0.62	2.8	0.97	1.17	0.95	1.31		
		THYMU1000037	1.53	4.15	2.11	2.06	2.81	1.45		
		THYMU1000042	5.97	10.24	12.23	12.03	13.98	13.28		
20		THYMU1000047	2.72	6.03	6.72	6.04	7.77	7.23		
		THYMU1000080	0.56	4.31	2.6	3.26	1.85	2.11		
		THYMU1000094	2.77	3.47	7.91	9.17	8.35	4.55		
25		THYMU1000109	17.28	14.34	111.37	98.05	142.29	93.04		
		THYMU1000127	2.75	5.95	10.76	8.18	9.98	6.74		
		THYMU1000130	2.5	4.4	4.55	6.69	6.07	4.94		
		THYMU1000137	3.53	7.18	10.26	12.67	18.55	13.05	*	+
30		THYMU1000146	4.37	8.38	6.52	8.29	7.46	7.74		
		THYMU1000159	5.43	9.51	16.37	12.4	15.15	13.27		
		THYMU1000163	5.85	12.26	37.58	45.53	58.37	36.93		
35		THYMU1000167	2.39	3.02	4.73	4.89	6.79	3.97		
		THYMU1000186	0.69	1.05	1.45	1.31	2.45	0.66		
		THYRO1000017	0.94	3.45	2.54	2.02	3.54	2.11		
		THYRO1000026	1.56	5.63	4.02	3.96	4.82	3.36		
40		THYRO1000034	0.49	4.16	1.59	1.99	2	1.82		
		THYRO1000035	0.86	4.84	1.34	2.29	2.48	2.11		
		THYRO1000036	0.93	8.32	4	3.08	4.36	5.59		
		THYRO1000040	2.58	7.02	4.76	4.66	4.83	4.93		
45		THYRO1000061	2.01	1.91	3.07	3.53	3.8	2.61		
		THYRO1000067	1.98	2.8	5.12	3.37	4.14	3.3		
		THYRO1000070	1.26	2.09	3.59	2.65	3.85	3.45		
50		THYRO1000072	1.33	3.37	4.22	2.54	4.08	2.06		
		THYRO1000084	8.07	12.69	22.39	2.99	5.61	4.42		
		THYRO1000085	1.44	5.66	3.99	2.42	3.86	2.85		
		THYRO1000086	-0.05	5.46	1.74	0.89	1.18	1.15		
55		THYRO1000087	0.72	3.86	1.01	0	0.58	0.17		

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	SKNMC1000004	9.7	11.62	16.77	10.19	12.16	13.96		
	SKNMC1000011	1.82	8.58	4.12	5.89	3.95	6.77		
5	SKNMC1000013	0.51	6.69	1.13	1.21	2.14	1.44		
	SKNMC1000014	1.28	4.18	3.22	3.77	6.37	3.96		
	SKNMC1000018	3.42	5.19	5.25	5.51	5.68	3.44		
	SKNMC1000020	0.95	4.03	3.46	3.6	4.68	4.56		
10	SKNMC1000046	2	3.17	3.48	3.95	3.26	2.55		
	SKNMC1000050	4.99	8.04	10.32	5.4	6.28	6.12		
	SKNMC1000062	9.79	12.6	20.18	19.2	15.42	18.73		
15	SKNMC1000075	1.45	4.3	2.01	1.98	1.89	2.92		
	SKNMC1000082	1.12	4.39	2.13	1.85	1.78	2.39		
	SKNMC1000091	4.54	7.52	7.95	11.74	12.86	12.77	**	+
	SKNMC1000099	0.33	4.29	1.98	1.32	0.65	1.18		
20	SKNMC1000104	1.13	4.24	3.45	1.47	3.14	2.43		
	SKNMC1000113	0.97	1.83	1.2	1.74	2.63	0.89		
	SKNMC1000119	1.73	2.64	5.07	4.48	5.34	4.67		
25	SKNMC1000142	0.04	2.87	0.99	1.27	0.75	1		
	SKNMC1000170	0.91	4.75	2.34	1.71	1.49	1.11		
	SKNMC1000178	3.02	8.39	7.08	5.77	9.65	9.02		
	SKNMC1000194	0.63	9.82	1.51	0.61	1.73	1.3		
30	SKNMC1000198	1.35	11.01	3.33	2.65	2.1	2.88		
	SKNMC1000225	1.35	6.44	2.97	2.39	3.4	3.26		
	SKNMC1000249	0.49	2.14	0.75	0.57	0.51	0.52		
35	SPLEN1000007	0.74	2.15	2.11	1.7	2.26	1.99		
	SPLEN1000012	0.39	1.9	1.72	1.19	0.8	0.84		
	SPLEN1000014	1.78	4.4	4.9	5.75	4.33	3.99		
	SPLEN1000036	4.95	11.64	24.32	20.56	27.73	21.68		
40	SPLEN1000059	0.04	6.69	1.06	0.91	1.79	1.47		
	SPLEN1000068	1.68	10.81	5.71	5.79	5.17	5.64		
	SPLEN1000072	1	8.5	4.7	2.82	3	2.21		
	SPLEN1000101	20.01	18.4	45.64	29.93	25.24	12.63		
45	SPLEN1000108	0.56	1.54	0.98	0.75	1.11	0.76		
	SPLEN1000113	1.33	2.27	3.04	2.72	4.13	3.04		
	SPLEN1000114	2.97	4.19	6.03	3.59	4.76	6.32		
50	SPLEN1000132	0.85	4	1.72	2.25	2.67	1.99		
	SPLEN1000135	3.13	8.76	14.93	11.12	15.28	10.52		
	SPLEN1000136	12.41	21.47	15.14	20.24	27.48	21.8		
	SPLEN1000141	2.26	7.07	10.79	4.03	5.41	4.51		
55	SPLEN1000164	2.49	3.79	8.58	3.98	5.88	7.61		

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	PLACE4000455	5.18	7.39	9.55	8	7.21	4.63		
	PLACE4000465	1.39	4.34	3.26	4.15	6.07	4.34		
5	PLACE4000466	120.96	98.04	201.25	113.83	170.96	145.31		
	PLACE4000472	3.12	9.6	10.17	10.92	13.21	9.22		
	PLACE4000487	3.18	7.83	16.5	14.66	16.62	15.05		
	PLACE4000489	0.93	4.69	3.41	1.95	3.88	1.69		
10	PLACE4000494	1.15	1.6	4.07	2.74	3.1	2.08		
	PLACE4000502	6.3	5.39	10.92	11.65	15.08	6.37		
	PLACE4000521	2.5	3.44	16.06	12.78	20.63	11.2		
15	PLACE4000522	5.07	6.17	9.07	12.43	8.68	14.11		
	PLACE4000537	0.98	4.28	1.27	1.67	1.61	1.22		
	PLACE4000548	1.99	5.69	2.46	3.04	3.68	2.32		
	PLACE4000558	0.87	6.72	1.97	3.15	2.41	2.15		
20	PLACE4000581	2.1	7.22	7.04	3.9	5.96	5.44		
	PLACE4000590	0.4	0.61	0.15	0.4	0.81	-0.25		
	PLACE4000593	2.94	2.98	5.22	4.44	5.82	3.83		
25	PLACE4000612	0.68	3.33	3.33	1.5	3.02	2.74		
	PLACE4000638	1.25	4.24	0.84	1.2	1.44	1.58		
	PLACE4000650	0.82	4.67	1.02	1.43	1.11	1.16		
	PLACE4000651	2.42	6.4	7.48	5	7.01	6.07		
30	PLACE4000654	0.98	5.7	2.47	1.35	2.48	1.47		
	PLACE4000670	0.5	4.06	2.92	0.76	1.29	0.67		
	PLACE4000685	6.35	8.68	13.83	13.46	14.26	13.77		
	PLACE4000687	0.37	3.02	1.11	2.2	1.4	1.12		
35	PLACE5000003	1.1	2.74	3.31	3.21	3.55	3.07		
	PLACE5000005	12.43	16.53	27.36	24.54	24.57	24.76		
	PLACE5000019	0.4	4.15	1.13	0.59	1.89	0.79		
40	PLACE5000021	0.74	4.59	1.61	0.39	0.93	0.32		
	PLACE5000022	1.2	6.11	2.25	3.17	2.76	2.09		
	PLACE5000024	1.77	2.58	2.27	2.92	3.39	3.84	*	+
	PLACE5000036	1.81	3.24	3.11	2.41	3.19	2.84		
45	PLACE5000059	14.41	17.79	26.55	25.98	30.03	34.87		
	PLACE5000076	1.41	3.61	2.22	4.04	3.96	2.54		
	PLACE5000117	7.44	12.48	15.66	16.87	18.78	20.64		
50	PLACE5000143	0.85	6.45	2.11	1.67	2.85	2.73		
	PLACE5000152	0.42	4.49	1.23	1.61	1.95	1.57		
	PLACE5000154	18.23	23.5	45.06	21.81	25.65	31.8		
	PLACE5000155	3.35	2.81	5.51	3.94	2.78	4.87		
55	PLACE5000165	3.78	4.4	6.67	4.51	5.99	5.82		

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	PLACE4000214	0.86	3.61	2.68	2.08	2.53	1.69		
	PLACE4000222	0.93	5.28	4.36	4.13	4.75	3.5		
5	PLACE4000223	0.46	4.51	1.79	1.37	1.22	0.38		
	PLACE4000229	1.9	5.79	2.11	2.81	3.36	3.48		
	PLACE4000230	1.11	5.89	6.51	3.61	6.81	5.15		
10	PLACE4000233	1.26	3.02	5.66	2.92	2.98	3.51		
	PLACE4000239	2.35	3.68	4.17	4.19	3.97	3.35		
	PLACE4000247	0.52	2.37	3.38	2.64	3.1	2.35		
	PLACE4000250	1.18	3.24	2.35	3.33	3.68	2.8		
15	PLACE4000252	1.06	4.99	2.25	1.92	1.75	1.64		
	PLACE4000259	4.42	11.95	18.1	14.47	22.09	14.02		
	PLACE4000261	0.87	10.29	1.07	2.03	1.9	1.12		
	PLACE4000264	15.86	24.96	36.9	11.96	21.82	22.51		
20	PLACE4000269	3.48	3.71	7.95	4.62	4.55	2.85		
	PLACE4000270	0.43	1.42	1.87	1.75	1.83	0.59		
	PLACE4000281	17.84	20.97	44.05	32.93	28.37	28.87		
25	PLACE4000300	0.67	2.06	2.04	3.21	2.88	3.58	*	+
	PLACE4000320	1.33	5.86	3.1	2.84	5.32	3.21		
	PLACE4000323	1.63	7.43	5.13	4.03	4.65	4.82		
	PLACE4000326	1.8	10.98	5.67	5.72	8.73	5.59		
30	PLACE4000344	0.22	5.75	2.62	1.66	1.6	1.18		
	PLACE4000347	4.7	3.82	13.93	16.83	16.75	17.36	*	+
	PLACE4000354	3.18	6.29	10.68	5.17	2.81	2.79		
	PLACE4000367	0.79	2.97	1.71	0.87	1.3	1.38		
35	PLACE4000369	1.35	3.97	2.36	1.99	1.96	0.82		
	PLACE4000379	2.44	6.66	5.44	5.94	7.55	5.07		
	PLACE4000387	0.88	5.86	2.11	1.28	0.84	1.12		
40	PLACE4000392	0.42	5.58	1.32	1.81	1.02	1.63		
	PLACE4000399	10.99	17.08	75.17	59.11	80.22	58		
	PLACE4000401	0.72	0.7	1.53	1.17	0.83	1.4		
	PLACE4000403	3.15	4.13	8.51	5.29	6.38	5.87		
45	PLACE4000411	2.22	2.28	4	2.27	2.6	1.82		
	PLACE4000415	0.7	3.55	2.8	1.16	1.86	0.78		
	PLACE4000416	25.49	29.13	33.54	23.65	21.92	24.83		
50	PLACE4000424	1.61	5.59	3.33	3.27	3.92	2.51		
	PLACE4000431	3.89	7.39	21.01	17.68	28.21	16.79		
	PLACE4000443	0.07	4.33	2.15	1.52	2.83	1.14		
	PLACE4000445	3.94	5.43	9.98	7.62	6.99	6.27		
55	PLACE4000450	2.99	3.65	23.28	15.51	24.53	16.04		

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	PLACE3000388	1.18	3.22	1.94	2.76	3.49	2.22		
	PLACE3000399	2.12	4.66	6.28	7.42	9.84	6.05		
5	PLACE3000400	3.08	5.44	11.87	7.97	10.77	7.82		
	PLACE3000401	7.52	11.42	18.59	22.61	29.55	23.4	*	+
	PLACE3000402	1.79	3.21	3.4	2.19	1.74	1.79		
	PLACE3000405	3.37	3.74	5.82	5.54	7.22	6.01		
10	PLACE3000406	2.1	2.91	3.11	3.48	3.68	2.42		
	PLACE3000413	1.18	2.72	2.69	1.71	2.06	1.52		
	PLACE3000416	1.05	4.03	4.04	3.43	2.67	2.72		
15	PLACE3000425	1.21	6.27	4.33	3.98	6.36	3.92		
	PLACE3000437	4.79	10.85	29.89	16.69	25.26	19.14		
	PLACE3000455	2.97	8.07	10.62	8.97	10.39	7.91		
	PLACE3000475	16.52	19.2	47.35	40.22	39.77	34.21		
20	PLACE3000477	5.44	4.79	5.56	8.05	5.52	8.42		
	PLACE4000003	0.38	2.97	1.61	3.14	2.33	2.31		
	PLACE4000008	15.19	11.38	16.76	13.05	14.26	8.84		
25	PLACE4000009	1.17	6.19	3.39	3.93	3.37	1.82		
	PLACE4000014	1.31	5.12	1.77	2.16	3.03	2.19		
	PLACE4000029	6.33	8.48	35.37	23.93	32.21	24.25		
	PLACE4000034	2.27	6.24	5.22	6.46	6.52	4.91		
30	PLACE4000049	3.39	3.35	5.21	3.85	5.82	4.86		
	PLACE4000052	1.41	3.36	2.2	2.62	2.64	2.02		
	PLACE4000062	1.6	4.94	5.06	4.25	5.06	3.71		
	PLACE4000063	2.59	6.87	5.19	4.86	4.81	3.73		
35	PLACE4000089	1.52	6.31	3.35	2.81	3.91	2.84		
	PLACE4000093	0.44	5.6	1.61	1.28	1.65	1.98		
	PLACE4000100	2.72	6.13	4.75	4.33	3.62	3.94		
40	PLACE4000103	0.63	4.48	5.64	4.4	5.67	2.9		
	PLACE4000106	3.2	5.33	6.63	7.1	5.13	7.21		
	PLACE4000128	1.93	3.97	4.88	4.15	4.96	4.4		
	PLACE4000129	0.74	3.26	1.64	1.57	2.11	1.78		
45	PLACE4000131	7.14	10.85	41.43	32.45	41.08	31.22		
	PLACE4000147	0.34	3.65	0.54	0.45	0.93	0.61		
	PLACE4000156	2.47	6.08	8.06	7.83	13.47	9.07		
	PLACE4000175	0.72	4.08	1.48	0.98	0.91	0.84		
50	PLACE4000190	14.55	18.47	70.34	49.15	74.82	60.76		
	PLACE4000192	1.3	2.27	3.6	2.36	2	1.25		
	PLACE4000206	5.35	6.65	12.44	7.13	7.1	6.02		
55	PLACE4000211	3.34	4.64	22.23	11.68	12.35	13.44		

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	PLACE3000199	0.22	4.8	1.63	0.55	0.86	0.43		
	PLACE3000205	11.79	17.95	57.49	51.4	63.83	42.63		
5	PLACE3000207	3.37	6.91	7.13	6.7	8.92	6.23		
	PLACE3000208	2.26	4.96	2.66	5.28	5.97	5.84		
	PLACE3000213	4.79	5.55	10.8	5.12	5.16	5.05		
	PLACE3000215	1.88	5.02	5.71	4.74	5	3.03		
10	PLACE3000218	0	1.63	1.18	0.97	0.62	0.31		
	PLACE3000220	1.96	3.55	4.58	6.74	7.52	6.69	*	+
	PLACE3000221	14.42	25.34	40.15	43.8	51.16	36.99		
15	PLACE3000225	1.15	4.68	3.11	2.11	2.67	1.28		
	PLACE3000226	1.37	5.65	5.16	3.78	7.42	3.68		
	PLACE3000230	0.83	3.46	1.36	2	2.8	1.73		
	PLACE3000231	1.31	1.97	2.37	4.86	3.95	4.12	**	+
20	PLACE3000235	1.12	1.75	3.89	3.95	4.21	3.39		
	PLACE3000242	2.6	5.11	9.24	9.46	10.97	8.29		
	PLACE3000244	1.05	3.2	1.81	1.85	1.81	0.64		
25	PLACE3000253	0.7	3.75	1.64	2.67	2.11	1.27		
	PLACE3000254	2.5	4.75	4.04	6.19	6.09	5.75	*	+
	PLACE3000271	2.67	6.06	6.81	10.96	10.99	9.5	*	+
	PLACE3000276	1.1	5.78	2.27	1.48	1.9	1.78		
30	PLACE3000304	5.55	4.69	10.81	11.19	11.49	10.5		
	PLACE3000309	0.43	1.67	1.87	2.43	2.94	2.78	*	+
	PLACE3000310	2.19	2.19	3.73	4.84	4.81	3.4		
	PLACE3000320	1.02	3.65	1.8	2.54	2.37	2.32		
35	PLACE3000322	1.31	4.23	6.63	7.5	7.8	6.09		
	PLACE3000330	24.05	24.44	41.08	31.87	35.83	29.17		
	PLACE3000331	1.21	5.86	4.14	4.34	5.7	4.31		
40	PLACE3000336	2.61	6.99	4.42	4.24	5.72	5.11		
	PLACE3000339	7.36	5.1	11.41	16.25	18.28	17.37	**	+
	PLACE3000341	1.65	1.32	2.41	4.08	4.35	3.65	**	+
	PLACE3000350	5.88	6.4	12.86	15.45	18.5	15.41	*	+
45	PLACE3000352	1.54	3.88	2.13	2.37	2.25	1.71		
	PLACE3000353	5.38	9.72	11.8	19.12	22.98	15.5	*	+
	PLACE3000362	0.62	4.92	4.72	3.61	5.33	3.39		
	PLACE3000363	2.19	5.13	2.32	1.89	3.28	2.07		
50	PLACE3000365	1.34	6.11	3.37	3.34	4.05	2.12		
	PLACE3000373	0.89	1.52	3.66	2.93	6.08	2.3		
	PLACE3000374	1.07	1.85	2.91	2.72	2.99	2.15		
55	PLACE3000387	0.31	3.32	1.04	1.24	1.65	1.29		

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	PLACE2000435	0.48	5.19	1.49	1.69	1.63	1.5		
	PLACE2000438	1.61	4.74	3.66	2.33	2.81	3.15		
5	PLACE2000450	3.01	4.38	5.67	6.51	7.39	5.63		
	PLACE2000455	0.24	2.62	1.24	1.65	2	1.82		
	PLACE2000458	0.38	3.3	1.81	1.06	2.7	1.24		
	PLACE2000464	2.15	4.91	5.3	7.43	9.68	8.83	*	+
10	PLACE2000465	1.43	6.72	6	6.51	8.27	6.31		
	PLACE2000473	120.94	179.35	328.3	214.7	297.75	279.74		
	PLACE2000477	0.43	3.87	1.34	1.13	2.22	0.97		
15	PLACE3000004	2.22	4.63	6.39	5.27	7.51	5.2		
	PLACE3000009	19.91	19.71	105.63	77.3	140.99	92.95		
	PLACE3000020	10.03	9.03	49.6	36.74	46.52	23.82		
	PLACE3000029	6.59	9.63	24.88	14.88	18.47	20.04		
20	PLACE3000038	0.52	2.37	2.47	1.44	2.4	2.05		
	PLACE3000052	5.13	7.95	23.92	25.01	29.61	24.94		
	PLACE3000059	0.57	5	2.42	0.75	2.8	1.27		
25	PLACE3000067	2.51	5.79	7.44	5.66	8.53	7.75		
	PLACE3000069	1.95	5.61	3.58	5.24	3.79	4.55		
	PLACE3000070	2.57	5.57	9.04	9.5	10.42	10.57		
	PLACE3000103	3.85	7.84	11.87	6.6	8.32	4.37		
30	PLACE3000119	1.59	2.74	3.15	3.24	3.67	2.95		
	PLACE3000121	7.58	8.44	38.1	30.63	42.28	32.64		
	PLACE3000124	1.53	4.54	5.95	6.35	7.75	7.18		
	PLACE3000135	0.69	5.46	1.1	0.76	0.9	0.59		
35	PLACE3000136	0.77	10.46	4.46	2.12	2.77	2.01		
	PLACE3000142	0.7	9.94	1.75	1.53	2.76	1.13		
	PLACE3000145	8.69	17.55	55.33	42.85	49.12	39.96		
40	PLACE3000147	15.7	12.92	39.97	64.96	54.76	30.34		
	PLACE3000148	0.7	2.08	1.48	0.82	1.4	1.38		
	PLACE3000154	0.48	1.86	0.67	1.5	0.87	0.44		
	PLACE3000155	1.28	4.26	2.53	3.97	4.76	3.04		
45	PLACE3000156	1.11	7.96	2.07	1.96	3.4	2.81		
	PLACE3000157	0.92	8.37	1.75	2.06	3.02	2.72		
	PLACE3000158	1.73	8.82	5.05	3.81	5.63	4.77		
50	PLACE3000160	8.11	15.22	19.5	33.66	34.71	33.78	**	+
	PLACE3000169	2.15	2.65	6.87	4.3	5.11	5.12		
	PLACE3000181	1.06	2.14	3.94	3.22	2.62	3		
	PLACE3000194	0.31	2.83	1.77	1.89	2.58	2.42		
55	PLACE3000197	1.18	2.66	2.09	2.42	2.46	2.21		

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	PLACE2000231	1.02	3.97	3.11	3.43	2.61	2.47		
	PLACE2000235	1.82	5.27	6.17	6.02	7.45	6.97		
5	PLACE2000246	1.93	2	6.06	4.58	5.09	3.93		
	PLACE2000264	0.67	1.39	1.85	2.45	3.74	3	*	+
	PLACE2000274	0.65	2.4	2.12	3.09	4.11	2.1		
10	PLACE2000287	0.81	4.44	1.49	2	2.59	1.34		
	PLACE2000296	1.01	4.56	1.55	2.5	3.16	1.69		
	PLACE2000302	1.34	4.67	2.86	3.52	3.35	3.45		
	PLACE2000305	3.09	6.65	5.42	6.3	7.15	5.02		
15	PLACE2000317	0.92	6.34	2.26	3.15	2.95	2.4		
	PLACE2000324	1.19	1.25	3.09	4.2	5.84	3.3		
	PLACE2000334	3.52	5	6.6	7.33	8.12	5.88		
20	PLACE2000335	1.47	4.35	4.24	5.68	6.25	4.76		
	PLACE2000340	0.64	3.47	1.63	1.58	1.52	1.65		
	PLACE2000341	4.21	7.87	28.81	18.94	32.45	19.16		
	PLACE2000342	2.07	5.11	4.32	4.84	5.82	4.49		
25	PLACE2000347	1.24	5.26	5.58	7.83	6.22	7.55		
	PLACE2000357	8.49	13.56	15.35	17.83	18.98	21.24	*	+
	PLACE2000358	2.87	3.65	8.67	4.88	7.63	4.37		
	PLACE2000359	1.27	1.79	4.45	3.28	6.65	3.61		
30	PLACE2000366	1.93	3.14	3.22	3.99	5.6	4.17		
	PLACE2000371	4.29	5.2	6.08	5.95	9.06	7.32		
	PLACE2000373	1.91	4.8	5.98	5.69	6.29	4.19		
	PLACE2000374	1.86	5.17	2.78	1.62	2.79	1.49		
35	PLACE2000379	0.34	4.85	1.32	1.28	0.92	0.04		
	PLACE2000386	39.29	43.92	84.66	87.53	104.55	76.56		
	PLACE2000388	1.96	3.35	3.89	2.78	3.48	3.49		
40	PLACE2000392	33.29	39.2	59.56	42.5	52.12	58.24		
	PLACE2000394	1.26	3.27	3.01	5.69	4.35	4.34	*	+
	PLACE2000398	0.73	3.88	2.36	2.03	3.59	1.35		
	PLACE2000399	3.7	6.82	7.01	7.15	7	6.79		
45	PLACE2000402	2.15	6.88	3.84	2.86	3.68	3.9		
	PLACE2000404	5.2	9.96	10.67	10.03	11.04	6.47		
	PLACE2000411	3.21	7.2	5.21	5.27	5.73	6.68		
50	PLACE2000418	0.73	2.28	2.41	2.22	3.07	2.37		
	PLACE2000419	0.99	2.32	2.54	4.95	4.55	3.29	*	+
	PLACE2000425	1.26	3.98	3.11	4.28	4.81	5.2		
	PLACE2000427	0.7	5.13	3.27	2.54	3.04	2.47		
55	PLACE2000433	0.77	7.05	2.6	2.33	3.09	2.46		

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	PLACE2000022	1.8	2.8	4.31	3.35	4.64	2.71
	PLACE2000030	9.37	11.08	71.38	55.43	60.32	35.97
5	PLACE2000032	1.23	2.89	3.51	3.53	3.24	2.32
	PLACE2000033	3.29	6.3	12.11	12.49	14.25	9.37
	PLACE2000034	0.6	4.3	1.92	1.79	2.19	2.44
	PLACE2000039	2.75	8.06	3.92	6.27	8.01	4.81
10	PLACE2000043	7	11.08	22.94	19.27	26.58	17.38
	PLACE2000044	0.63	6.92	1.12	1.8	1.71	1.29
	PLACE2000047	0.84	5.25	5.15	5.4	6.46	7.32
15	PLACE2000050	1.48	2.68	4.78	2.92	2.98	1.2
	PLACE2000061	0.47	2.07	1.17	0.95	1.34	0.25
	PLACE2000062	1.99	2.83	4.12	4.89	4.4	3.09
	PLACE2000072	0.78	2.45	1.57	1.85	1.62	1.69
20	PLACE2000073	0.89	5.86	2.86	2.8	2.61	3.07
	PLACE2000097	8.54	19.93	23.93	27.69	35.36	26.36
	PLACE2000100	1.87	7.79	3.53	4.23	6.29	4.22
25	PLACE2000103	1.03	7.44	2.25	3.51	5.17	4.38
	PLACE2000106	1.53	2.42	4.19	5.71	3.29	4.01
	PLACE2000111	2.05	3.17	4.37	4.07	6.6	4.75
	PLACE2000115	0.3	2.06	0.75	0.31	1.29	0.97
30	PLACE2000118	10.15	17.04	21.09	13.73	15.21	22.01
	PLACE2000124	10.14	17.83	62.13	53.2	98.37	62.96
	PLACE2000132	0.06	6.26	1.48	0.99	1.5	1.56
35	PLACE2000136	0.55	7.94	0.91	1.47	1.37	1.02
	PLACE2000137	0.96	4.46	2.4	2.65	4.12	3.14
	PLACE2000140	2.91	5.24	14.34	13.2	12.08	9.43
	PLACE2000147	1.49	1.52	2.83	1.06	0.97	1.12
40	PLACE2000153	0.3	3.44	2.15	1.69	2.45	1.95
	PLACE2000164	0.66	2.78	1.13	2.66	1.62	1.62
	PLACE2000170	1.54	6.18	4.69	5.26	9.09	6.24
	PLACE2000172	0.33	4.34	2.15	1.25	1.93	2.43
45	PLACE2000173	0.92	4.97	3.37	3.33	3.71	2.74
	PLACE2000174	1.17	4.68	2.43	1.85	2.89	2.05
	PLACE2000176	1.22	1.57	2.58	3.28	2.27	1.42
50	PLACE2000187	1.01	2.08	2.55	3.45	3.66	2.19
	PLACE2000216	7.03	9.28	11.47	14.09	9.13	3.68
	PLACE2000219	0.69	4.02	2.72	3.55	3.58	2.3
	PLACE2000221	2.49	6.81	6.53	7.22	9.33	8.56
55	PLACE2000223	0.72	3.2	1.71	1.16	1.05	0.48

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	PLACE1011795	0.74	1.28	2.24	1.47	1.8	1.22		
	PLACE1011810	9.28	10.82	19.51	13.32	17.73	15.47		
5	PLACE1011824	5.38	8.17	38.05	25.52	42.89	16.75		
	PLACE1011825	10.61	16.39	22.6	17.63	18.92	16.07		
	PLACE1011835	24.64	32.67	47.67	32.09	26.75	37.23		
	PLACE1011836	18.11	18.97	31.43	33.14	47.23	13.95		
10	PLACE1011847	2.67	6.74	6.42	5.34	7.18	5.84		
	PLACE1011855	0.9	6.06	3.53	3.23	4.47	3.49		
	PLACE1011858	5.83	7.44	9.37	6.88	6.25	8.35		
	PLACE1011874	1.35	3.14	3.55	4.18	5.28	4.65		
15	PLACE1011875	0.57	2.29	1.11	2.66	2.48	2.64		
	PLACE1011877	3.8	5.03	4.4	7.67	6.8	9.97	*	+
	PLACE1011891	0.17	3.81	1.31	1.13	1.34	1.26		
20	PLACE1011896	-0.1	5.22	1.45	-0.19	1.07	0.32		
	PLACE1011920	0.21	5.87	1.04	1.1	1.92	1.76		
	PLACE1011922	2.4	4.18	4.72	3.72	2.88	3.98		
	PLACE1011923	3.42	4.82	7.51	8.09	5.38	12.28		
25	PLACE1011937	3.16	2.24	3.76	3.81	5.58	4.64		
	PLACE1011939	14.93	17.81	26.01	41.75	45.05	47.88	**	+
	PLACE1011940	6.13	7.58	12.68	5.73	7	7.23		
30	PLACE1011962	3.28	7.83	7.35	6.39	8.38	8.01		
	PLACE1011964	0.39	5.05	1.66	1.04	1.95	1.46		
	PLACE1011978	1.55	4.65	3.35	4.48	5.91	5.14		
	PLACE1011980	2.1	4.62	6.07	3.95	4.91	5.35		
35	PLACE1011981	5.77	7.28	38.2	27.22	35.24	34.99		
	PLACE1011982	0.83	3.1	2.23	1.82	2.59	1.57		
	PLACE1011995	0.81	3.4	3.73	2.51	3.59	2.86		
40	PLACE1012023	1.38	5.37	1.87	2.09	2.46	1.91		
	PLACE1012026	1.95	5.72	4.23	6.08	9.51	9.8		
	PLACE1012031	2.49	5.81	4.54	4.34	6.35	4.03		
	PLACE2000003	1.18	3.64	6.86	7.38	8.12	8.92		
45	PLACE2000005	1.16	2.41	2.16	2.76	2.03	1.89		
	PLACE2000006	2.52	4.13	15.6	11.34	16.58	13.4		
	PLACE2000007	0.96	4.85	4.24	3.94	5.13	3.33		
	PLACE2000011	1.72	3.27	3.34	4.3	5.06	3.62		
50	PLACE2000014	4.04	5.93	23.94	27.19	30.87	30.97	*	+
	PLACE2000015	1.27	4.79	3.52	2.77	3.31	2.69		
	PLACE2000017	0.48	4.78	2.15	2.65	2.56	2.45		
55	PLACE2000021	1.99	5.06	4.09	3.72	6.24	5.61		

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	PLACE1011478	1.24	1.64	3.11	5.03	3.02	3.01		
	PLACE1011492	2.24	3.54	5.01	5.45	6.74	5.27		
5	PLACE1011498	0.57	2.77	0.98	1.18	1.69	1.15		
	PLACE1011501	0.49	4.67	1.8	4.87	5.63	7.5		
	PLACE1011503	0.44	5.37	0.79	0.5	0.82	0.35		
10	PLACE1011509	1.1	7.8	3.38	4.62	5.08	5.74		
	PLACE1011514	5.86	11.61	11.98	13	17.7	13.17		
	PLACE1011516	10.37	13.29	18.32	18.32	8.08	3.63		
	PLACE1011520	0.34	0.95	1.34	1.3	1.73	1.01		
15	PLACE1011538	52.87	99.27	185.04	129.45	86.87	63.9		
	PLACE1011555	0.87	2.88	2.83	1.55	2.03	1.48		
	PLACE1011561	3.2	4.53	1.98	6.81	5.31	3.08		
	PLACE1011563	1.35	4.74	3.49	2.52	2.64	2.87		
20	PLACE1011567	1.04	4.94	3.36	2.75	4.19	2.77		
	PLACE1011569	0.32	4.35	2.77	2.38	2.46	2.28		
	PLACE1011576	3.25	1.88	7.94	7.85	9.1	8.03		
25	PLACE1011586	3.24	2.5	6.22	4.43	4.35	2.94		
	PLACE1011635	1.85	2.56	4.53	9.96	12.43	8.71	**	+
	PLACE1011641	0.43	2.9	0.9	1.71	1.18	1.19		
	PLACE1011642	5.05	6.96	10.37	12.13	10.86	10.09		
30	PLACE1011643	1.29	3.69	1.14	2.38	2.28	2.14		
	PLACE1011646	8.68	12.8	30.53	39.05	54.16	37.76	*	+
	PLACE1011649	1.35	5.7	3.88	4.5	4.61	5.46		
35	PLACE1011650	1.15	1.45	2.54	2.47	3.24	3.75		
	PLACE1011661	1.02	2.26	2.8	3.95	5.92	3.46		
	PLACE1011664	2.21	3.18	3.99	5.31	3.93	1.73		
	PLACE1011672	0.88	4.14	0.72	2.69	3.57	2.04		
40	PLACE1011675	0.51	2.31	1.32	1.66	0.99	0.83		
	PLACE1011682	2.04	4.56	2.23	2.22	4.03	2.12		
	PLACE1011708	1.1	5.89	3.8	5.5	8.12	4.66		
	PLACE1011719	1.07	4.58	1.66	3.55	3.88	3.03		
45	PLACE1011725	1.23	1.19	2.72	3.73	5.19	4.3	*	+
	PLACE1011729	0.86	1.03	1.8	2.38	3.26	1.22		
	PLACE1011741	2.36	3.67	3.64	4.16	2.52	4.23		
50	PLACE1011749	1.58	3.89	4.09	3.49	4.85	3.27		
	PLACE1011757	20.92	30.53	55.88	56.6	55.88	49.59		
	PLACE1011762	0.4	4.34	2.69	3.91	2.14	2.3		
	PLACE1011778	0.51	4.39	1.99	1.34	2.02	1.34		
55	PLACE1011783	2.59	4.63	5.46	4.8	8.41	5.55		

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	PLACE1011185	2.47	4	3.57	4.66	5.15	4.8	*	+
	PLACE1011186	13.16	12.8	16.45	21.18	25.69	28.83	*	+
5	PLACE1011203	1.08	4.64	2.19	1.75	2.94	1.96		
	PLACE1011214	9.02	16.55	59.24	46.39	62.58	47.89		
	PLACE1011219	1.41	4.91	3.6	3.26	4.96	2.9		
10	PLACE1011221	2.68	5.47	6.15	6.23	8.57	7.51		
	PLACE1011229	1.38	4.2	2.69	2.43	2.43	3.09		
	PLACE1011231	0.53	1.5	1.62	1.84	2.59	2.4		
	PLACE1011236	5.69	7.9	43.53	33.48	56.77	44.25		
15	PLACE1011247	8.36	10.08	19.16	24.17	26.33	29.74	*	+
	PLACE1011263	0.57	6.43	4.23	2.62	4.11	5.09		
	PLACE1011273	0.72	3.62	1.17	2.01	1.64	1.72		
20	PLACE1011278	2.42	5	6.12	3.98	4.84	4.31		
	PLACE1011289	2.73	5.84	7.57	6.34	6.13	6.08		
	PLACE1011291	3.2	5.19	8.31	7.32	8.04	7.95		
	PLACE1011296	0.93	2.45	1.94	1.76	2.63	2.05		
25	PLACE1011310	1.72	2.64	3.36	4.51	2.77	4.48		
	PLACE1011311	1.8	3.97	6.33	7.8	9.34	6.73		
	PLACE1011321	1.29	4.77	3.5	3.3	2.63	3.33		
	PLACE1011325	0.63	4.2	1.84	2	2.74	1.59		
30	PLACE1011332	5.65	10.55	9.4	14.8	14.57	15.04	*	+
	PLACE1011340	0.86	4.88	3.38	3.81	4.71	3.47		
	PLACE1011353	5.39	5.53	8.39	10.06	8.58	4.43		
35	PLACE1011360	1.09	3.29	2.18	2.11	3.06	2.41		
	PLACE1011364	4.88	5.69	7.92	6.34	4.57	4.57		
	PLACE1011365	0.92	3.36	2.95	3.01	3.6	2.06		
	PLACE1011371	9.37	10.69	63.06	56.51	87.51	64.33		
40	PLACE1011375	1.62	9.37	3.35	2.61	2.74	1.62		
	PLACE1011386	3.91	12.96	4.18	6.88	5.51	5.93		
	PLACE1011399	1.52	10.14	4.27	2.12	4.22	3.77		
45	PLACE1011406	1.25	2.05	1.81	5.03	3.26	2.38		
	PLACE1011407	0.64	2.05	2.58	2.08	2.19	2.02		
	PLACE1011419	2.83	3.2	6.02	6.23	4.18	3.72		
	PLACE1011433	3.09	4.43	6.13	3.41	4.74	4.99		
50	PLACE1011440	0.77	5.81	2.48	2.1	2.24	2.27		
	PLACE1011452	1.06	8.86	2.43	2.87	2.71	2.68		
	PLACE1011465	0.09	8.53	1.96	1.04	1.68	0.85		
	PLACE1011472	1.52	7.67	3.98	0.97	2.87	1.59		
55	PLACE1011477	11.18	12.29	54.35	31.86	55.86	45.67		

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	PLACE1010924	1.15	2.31	1.55	1.11	1.88	0.92		
	PLACE1010925	2.76	5.36	2.17	4.92	6.32	4.16		
5	PLACE1010926	1.8	5.73	4.31	5.37	4.35	4.45		
	PLACE1010942	1.7	6.25	5.63	5.53	7.88	7.69		
	PLACE1010943	7.38	10.43	17.12	24.62	29.5	31.96	**	+
10	PLACE1010944	4.33	7.39	9.3	13.11	11.44	15.58	*	+
	PLACE1010947	1.43	0.9	2.41	2.57	2.8	2.25		
	PLACE1010954	3.56	2.92	7.4	7.92	8.55	7.64		
	PLACE1010960	2.06	3.44	6.07	4.23	7.89	2.6		
15	PLACE1010965	2.33	3.81	3.54	6.09	4.92	5.03	*	+
	PLACE1010968	1.55	4.69	1.38	4.95	6.51	3.68		
	PLACE1010978	3.63	6.12	7.05	9	8.94	6.45		
20	PLACE1010982	2.23	5.77	5.6	4.74	5.66	4.88		
	PLACE1010990	0.88	5.4	2.04	3.11	3.03	2.19		
	PLACE1011017	5.6	3.78	22.57	25.64	35.47	20.97		
	PLACE1011019	1.1	1.5	2.48	3.78	4.42	0.88		
25	PLACE1011026	4.17	4.93	6.23	6	5.19	2.74		
	PLACE1011032	0.89	3.95	2.03	1.44	1.58	0.56		
	PLACE1011041	1.07	4.13	1.03	1.69	1.22	1.43		
30	PLACE1011045	1.49	5.62	2.36	3.26	3.67	4.45		
	PLACE1011046	0.83	5.25	1.79	2.57	3.17	1.87		
	PLACE1011054	2.33	5.33	6.64	5.26	7.29	5.05		
	PLACE1011056	5.78	5.43	16.22	14.56	19.78	15.67		
35	PLACE1011057	2	2.18	3.5	3.29	5.68	3.9		
	PLACE1011059	0.93	1.37	1.56	1.74	2.96	1.79		
	PLACE1011066	4.49	5.74	6.76	5.38	7.72	5.49		
	PLACE1011087	7.6	7.04	16.48	12.43	17.42	9.79		
40	PLACE1011090	2.98	6.14	6.74	3.36	4.13	3.26		
	PLACE1011109	1.99	7.29	4.29	4.08	7.96	3.83		
	PLACE1011114	1.62	4.4	3.13	3.33	4.68	3.29		
45	PLACE1011116	4.89	5.94	6.66	7.43	6.81	8.98		
	PLACE1011122	0.93	2.52	1.84	2.1	1.61	1.64		
	PLACE1011133	0.83	2.22	3.03	3.48	3.52	2.77		
	PLACE1011134	12.47	15.29	66.86	44.95	68.68	51.65		
50	PLACE1011143	0.68	4.53	1.48	1.84	2.62	1.41		
	PLACE1011146	0.91	5.93	1.74	1.97	3.23	2.36		
	PLACE1011160	1.67	7.36	3.81	3.42	4.53	4.24		
	PLACE1011165	1.77	2.34	3.39	2.15	3.39	3.8		
55	PLACE1011181	5.25	8.31	37.21	29.38	38.44	30.55		

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	PLACE1010629	1.08	4.64	2.24	3.01	3.3	3.5		
	PLACE1010630	1.64	4.77	3.69	4.22	4.78	5.29		
5	PLACE1010631	0.5	5.35	2.64	1.55	1.66	2.67		
	PLACE1010651	14.24	15.75	24.44	37.62	40.09	52.12	*	+
	PLACE1010661	1.62	4.09	2.28	3.56	6.43	3.22		
10	PLACE1010662	1.32	2.48	1.6	2.98	2.3	1.61		
	PLACE1010668	12.87	15.91	27.82	37.63	30.53	28.75		
	PLACE1010702	1.46	2.34	4.24	3.59	3.6	3.2		
	PLACE1010709	79.16	78.33	115.91	107.07	96.3	133.25		
15	PLACE1010713	7	10.81	14.7	9.14	8.16	15.14		
	PLACE1010714	0.82	7.41	1.58	1.75	2.04	1.47		
	PLACE1010716	0.71	6.19	4.31	2.08	2.3	1.93		
	PLACE1010717	0.9	6.49	2.13	2.17	3.9	2.61		
20	PLACE1010720	14.03	17.05	53.79	46.72	50.7	41.49		
	PLACE1010739	0.9	1.2	1.11	1.73	1.21	1.93		
	PLACE1010743	1.09	2.3	1.99	2.63	2.05	2.21		
25	PLACE1010752	0.87	2.92	1.85	1.76	2.05	1.53		
	PLACE1010761	3.6	8.83	13.51	12.07	16.4	17.08		
	PLACE1010771	1.41	6.89	5.03	6.13	10.3	5.42		
	PLACE1010784	0.9	9.66	1.97	2.07	1.72	1.34		
30	PLACE1010786	1.21	8.77	2.83	2.91	3.68	2.21		
	PLACE1010789	0.6	1.16	1.52	1.8	1.89	1.17		
	PLACE1010800	2.18	2.86	3.25	3.95	3.24	2.93		
35	PLACE1010802	2.97	4.63	5.31	5.72	4.27	3.3		
	PLACE1010811	0.89	2.19	1.96	1.83	1.75	2.18		
	PLACE1010813	8.89	13.3	55.85	48.82	72.26	46.7		
	PLACE1010827	1.54	6.43	3.94	4.3	5.52	4.81		
40	PLACE1010833	0.93	8.13	2.63	2.68	3.64	2.09		
	PLACE1010839	1.57	6.22	3.21	4.22	6.72	4.13		
	PLACE1010856	7.58	8.94	12.34	8.02	6.53	8.59		
	PLACE1010857	3.41	3.81	7.63	8.24	5.98	4.56		
45	PLACE1010870	1.3	2.24	2.05	1.62	2.38	1.94		
	PLACE1010877	1.67	4.66	2.69	2.77	3.92	2.62		
	PLACE1010882	0.49	4.8	0.99	1.74	1.27	0.51		
50	PLACE1010891	1.1	7.73	1.12	1.85	1.73	0.95		
	PLACE1010896	1.19	5.29	3.18	3.98	3.05	3.47		
	PLACE1010900	7.41	13.29	27.9	23.88	20.99	18.85		
	PLACE1010916	1.55	1.18	2.06	1.89	1.9	2.2		
55	PLACE1010917	-0.04	0.82	0.36	1.05	2.25	0.56		

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	PLACE1010324	0.88	3.49	1.56	1.12	1.54	1.2		
	PLACE1010329	0.73	4.64	1.95	2.56	3.37	1.96		
5	PLACE1010330	3.78	9.09	7.29	2.42	10.45	7.67		
	PLACE1010335	6.43	11.15	7.43	13.15	17.89	19.1	*	+
	PLACE1010341	0.19	4.81	1.24	1.07	3.13	1.54		
10	PLACE1010342	0.77	0.9	0.75	0.48	2.12	0.8		
	PLACE1010346	1.47	1.73	3.61	2.96	4.47	1.71		
	PLACE1010362	1.31	2.69	2.22	2.18	3.49	3.31		
	PLACE1010364	0.78	2.56	1.65	1.19	2.32	1.49		
15	PLACE1010368	1.66	5.44	3.51	3.41	3.87	4.48		
	PLACE1010373	9.05	10.48	16.82	12.13	15.45	12.28		
	PLACE1010383	1.91	5.52	5.13	5.58	6.39	4.9		
20	PLACE1010385	0.3	3.01	1.07	0.04	0.6	0.9		
	PLACE1010389	6.28	7.98	13.24	22.3	13.64	22.94	*	+
	PLACE1010401	0.73	2.72	1.32	1.99	2.87	2.21		
	PLACE1010410	3.15	4.83	6.71	4.78	7.55	7.32		
25	PLACE1010418	1.88	4.73	4.2	4.71	5.14	4.66		
	PLACE1010425	0.93	4.78	1.43	1.78	1.96	2.15		
	PLACE1010443	6.98	13.83	51.39	36.22	63.67	48.93		
	PLACE1010445	0.95	5.02	0.68	2.69	3.81	2.44		
30	PLACE1010481	1.19	2.06	2.46	1.75	2	1.85		
	PLACE1010482	28.99	29.39	53.06	31.75	19.63	40.44		
	PLACE1010491	3.36	6.4	5.38	4.96	5.98	3.26		
35	PLACE1010492	4	4.75	5.95	5.57	8.15	7.77		
	PLACE1010509	0.8	4.32	3.15	3.09	3.01	2.99		
	PLACE1010518	3.33	6.72	6.58	7.3	8.25	8.24		
	PLACE1010522	2.3	5.96	4.35	1.87	3.52	2.64		
40	PLACE1010529	1.8	6.5	3.84	5.43	8.44	7.2		
	PLACE1010547	0.79	1.57	1.76	1.51	1.93	2.45		
	PLACE1010560	0.63	2.51	2.06	3.31	2.2	2.51		
45	PLACE1010562	0.74	2.68	1.65	1.64	2.1	1.63		
	PLACE1010579	1.11	7.13	2.63	3.45	4.67	3.26		
	PLACE1010580	1.35	9.12	3.79	3.27	2.49	4.52		
	PLACE1010599	3.56	6.07	7.94	8.32	9.26	8.13		
50	PLACE1010606	1.17	4.42	1.64	3.39	4.17	3.51		
	PLACE1010616	1.84	3.72	5.49	4.09	8.09	6.81		
	PLACE1010622	2.1	3.43	3.91	5.07	4.74	4.69	*	+
	PLACE1010624	1.43	3.35	3.98	4.21	6.17	5.22		
55	PLACE1010628	1.37	3.97	2.83	2.14	3.63	4.22		

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	PLACE1010060	1.63	6.1	4.13	4.11	4.6	4.05		
	PLACE1010069	0.41	7.96	2.32	1.48	2.91	1.3		
5	PLACE1010070	0.92	8.04	1.5	0.45	1.78	1.09		
	PLACE1010074	5.25	11.67	11.8	12.32	9.51	11.22		
	PLACE1010076	12.75	11.95	29.01	19.58	15.88	16.82		
	PLACE1010078	2.96	2.42	4.36	4.64	4.39	3.85		
10	PLACE1010081	2.74	4.1	3.7	6.85	7.81	4.59	*	+
	PLACE1010083	0.69	2.53	1.51	1.22	1.86	2.26		
	PLACE1010089	1.86	4.89	3.35	3.45	3.47	3.48		
15	PLACE1010096	2.17	7.73	2.91	3.43	4.19	3.2		
	PLACE1010102	3.89	10.9	5.33	7.1	9.64	7.57		
	PLACE1010105	2.98	7.46	4.93	7.31	9.04	9.82		
	PLACE1010106	2.46	2.95	4.48	5.47	4.88	5.8	*	+
20	PLACE1010130	0.53	1.79	1.17	2.27	2.55	1.23		
	PLACE1010132	2.49	4.65	5.3	5.07	4.39	4.19		
	PLACE1010134	0.8	3.32	1.15	1.97	2.31	1.67		
25	PLACE1010139	6.67	10.51	12.98	14.99	16.1	14.58		
	PLACE1010148	0.96	5.07	1.62	1.48	1.9	1.97		
	PLACE1010152	3.11	5.68	5.16	6.33	6.64	5.76		
	PLACE1010155	3.8	6.17	6.52	16.85	20.56	20.32	**	+
30	PLACE1010156	13.71	15.43	32.21	132.45	85.59	134.99	**	+
	PLACE1010161	1.9	2.81	5.05	3.29	2.92	1.97		
	PLACE1010181	0.73	2.22	1.51	2.58	1.99	2.53		
	PLACE1010194	0.64	3.35	1.03	2.26	2.14	1.64		
35	PLACE1010202	0.4	4.14	1.2	2.91	1.65	2.16		
	PLACE1010231	1.1	3.78	2.39	1.31	2.99	2.73		
	PLACE1010235	1.26	4.24	1.94	2.68	2.42	3.16		
40	PLACE1010237	1.01	3.4	2.1	1.14	1.97	0.87		
	PLACE1010251	0.59	0.98	1.95	2.57	3.18	1.62		
	PLACE1010261	0.97	2.63	2.07	2.69	1.69	1.55		
	PLACE1010270	0.76	2.7	1.3	1.39	2.33	1.42		
45	PLACE1010273	0.97	3.27	0.46	1.48	2.25	1.5		
	PLACE1010274	6.28	9.23	9.66	10.49	12.18	14.28		
	PLACE1010277	6.03	10.14	12.68	14.6	16.06	15.84	*	+
	PLACE1010293	1.8	5.68	3.55	3.65	3.37	3.96		
50	PLACE1010297	5.17	11.37	21	24.84	32.59	22.06		
	PLACE1010300	4.18	4.78	8.22	8	9.95	6.87		
	PLACE1010310	16.52	14.75	49.45	70.74	71	77.01	*	+
55	PLACE1010321	2.03	4.92	2.46	3.37	4.99	2.73		

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	PLACE1009763	3.95	9.73	6.82	7.13	7.44	8.39
	PLACE1009766	1.46	6.98	3.32	3.07	5.19	3.75
5	PLACE1009772	0.48	5.19	0.6	1.01	2.46	0.89
	PLACE1009782	0.91	2.39	2.03	2.88	2.91	3.74
	PLACE1009794	2.58	4.45	5.11	3.54	3.66	5.03
	PLACE1009798	1.59	5.37	4	6.26	5.57	5.67
10	PLACE1009845	1.05	6.02	2.92	2.79	3.39	3.92
	PLACE1009849	0.96	6.61	2.35	1.79	3.41	2.59
	PLACE1009857	0.79	4.86	1.45	1.19	1.27	1.56
15	PLACE1009861	1.43	4.67	3.87	4.1	3.47	3.11
	PLACE1009872	53.53	52.43	88.5	74.95	49.47	81.73
	PLACE1009877	5.45	7.59	12.08	10.03	10.3	12.32
	PLACE1009879	0.82	3.28	1.59	1.55	2.99	1.34
20	PLACE1009886	0.68	4.04	1.53	1.62	3.04	1.72
	PLACE1009888	1.03	7.4	3.23	5.34	5.84	7.94
	PLACE1009908	1.56	7.63	8.64	3.37	6.71	5.9
25	PLACE1009919	4.5	7.53	8.26	5.84	10.72	10.15
	PLACE1009921	0.96	3.94	3.32	1.63	4.28	2.47
	PLACE1009923	3.82	5.56	6.85	6.32	8.13	5.57
	PLACE1009924	3.01	2.49	4.53	4.43	4.31	1.04
30	PLACE1009925	0.61	2.77	1.84	2.51	2.2	2.5
	PLACE1009931	2.78	5.21	9	8.71	6.93	8.09
	PLACE1009935	0.74	3.71	2.1	1.19	1.08	1.5
35	PLACE1009947	0.47	3.83	1.64	1.51	2.46	2.03
	PLACE1009961	0.43	4.08	1.39	1.69	2.18	1.9
	PLACE1009971	0.92	4.9	1.98	1.35	1.74	1.45
	PLACE1009982	40.34	48.71	89.8	53.8	57.87	66.96
40	PLACE1009992	0.94	1.9	2.59	1.47	2.52	0.68
	PLACE1009995	6.47	10.83	15.72	7.79	9.03	11.23
	PLACE1009997	0.55	3.7	3.03	2.76	3.2	2.64
45	PLACE1010002	1.4	4.14	2.82	2.89	3.04	3.46
	PLACE1010011	2.09	8.13	3.85	4.4	5.21	4.68
	PLACE1010013	0.18	12.85	1.74	0.92	1.81	0.68
	PLACE1010021	3.18	11.98	5.42	4.12	4.13	6.06
50	PLACE1010023	2.15	8.45	5.16	5.62	6.52	6.14
	PLACE1010031	4.6	4.35	7.23	6.79	4.91	1.82
	PLACE1010039	1.17	3.3	1.45	1.28	1.23	1.19
	PLACE1010045	1.1	2.83	3.66	2.52	3.55	2.64
55	PLACE1010053	1.42	3.56	1.65	2.21	2.76	2.37

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	PLACE1009527	0.95	4.51	2.11	1.64	2.55	1.28		
	PLACE1009531	20.82	28.24	41.52	46.25	43.25	49.96		
5	PLACE1009535	1.1	1.56	2.68	2.42	2.15	1.11		
	PLACE1009539	2.15	3.41	4.18	3.88	2.65	2.57		
	PLACE1009540	5.89	8	11.66	14.8	4.47	3.84		
10	PLACE1009542	1.11	3.37	1.42	1.51	2.06	1.44		
	PLACE1009546	0.62	5.27	0.97	2.24	1.64	1.25		
	PLACE1009556	0.35	4.46	3.46	3.36	2.86	3.16		
	PLACE1009569	0.05	3.93	2.46	3.34	2.7	3.11		
15	PLACE1009571	1.67	4.27	2.52	3.04	2.85	2.67		
	PLACE1009573	3.81	2.97	6.73	6.92	8.12	6.49		
	PLACE1009576	1.92	2.51	4.3	2.73	3.66	2.08		
20	PLACE1009580	1.42	1.81	1.74	2.73	3.47	2.33	*	+
	PLACE1009581	0.89	4.25	2.03	2.91	4.38	2.74		
	PLACE1009587	0.96	4.91	2.29	2.43	3.2	1.99		
	PLACE1009593	2.71	6.73	4.37	4.94	6.85	5.03		
25	PLACE1009595	1.81	5.44	2.66	2.67	5	2.79		
	PLACE1009596	1.57	6.83	2.6	3.44	3.97	2.7		
	PLACE1009600	3.03	4.27	4.48	5.48	9.14	4.42		
	PLACE1009604	2.32	4.64	5.02	4.22	6.11	3.23		
30	PLACE1009607	1.29	2.48	3.18	3.19	4.17	3.18		
	PLACE1009613	1.94	5.23	2.94	2.65	3.08	2.23		
	PLACE1009621	1.66	6.72	3.32	8.21	8.67	8.06	*	+
35	PLACE1009622	1.78	5.93	3.78	3.9	4.1	3.9		
	PLACE1009624	1.16	5.77	3.42	3.2	3.65	3.5		
	PLACE1009637	2	6.88	3.36	3.07	4.59	3.91		
	PLACE1009639	1.94	1.76	4.15	3.44	3.67	4.99		
40	PLACE1009654	20.88	17.13	34.95	14.94	24.53	20.64		
	PLACE1009659	2.77	6.78	7.45	6.38	8.38	6.55		
	PLACE1009665	1.04	4.21	1.93	1.19	2.72	1.93		
	PLACE1009669	7.73	9.64	14.54	9.85	16.89	8.82		
45	PLACE1009670	1.76	5.36	2.54	2.77	4.47	4.01		
	PLACE1009708	2.1	5.57	5.09	3.64	6.54	5.84		
	PLACE1009721	1.34	4.28	3.56	5.78	5.81	3.01		
50	PLACE1009731	1.36	3.59	3	3.58	6.53	5		
	PLACE1009735	1.94	3.94	3.21	5.16	7.52	4.78		
	PLACE1009737	1.89	4.29	2.95	4.83	5.61	5.47	*	+
	PLACE1009741	1.3	4.32	3.45	2.09	5.03	3.07		
55	PLACE1009752	1.34	5.64	2.65	2.3	3.33	1.68		

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	PLACE1009174	1.13	3.45	2.42	1.67	3.02	2.38		
	PLACE1009183	1.62	3.54	4.47	4.1	6.33	8.06		
5	PLACE1009186	1.04	5.07	2.3	2.46	2.86	2.91		
	PLACE1009190	0.75	2.32	1.44	1.53	1.9	1.65		
	PLACE1009196	0.81	4.01	2.73	2.24	2.38	1.99		
10	PLACE1009200	1.01	4.44	2.94	2.84	4.39	2.91		
	PLACE1009217	2.55	4.91	3.43	4.46	7.29	7.23		
	PLACE1009230	1.9	5.55	6.63	5.63	9.16	9.46		
	PLACE1009236	4.97	7.07	12.6	8.21	10.79	7.13		
15	PLACE1009246	11.71	11.96	24.75	14.59	16.36	9.05		
	PLACE1009265	6.95	7.82	14.01	15.61	5.19	12.17		
	PLACE1009279	0.67	2.07	2.46	1.93	2.54	1.63		
20	PLACE1009298	5.54	9.92	9.52	10.21	11.25	17.55		
	PLACE1009308	1.13	6.82	2.04	2.48	2.48	2.34		
	PLACE1009319	2.04	9.25	3.15	2.92	3.54	2.5		
	PLACE1009328	1.04	5.78	1.81	2.98	3.39	2.17		
25	PLACE1009335	1.38	6.55	4.72	2.24	3.21	3.01		
	PLACE1009338	2.56	4.14	5.1	3.24	4.3	1.57		
	PLACE1009344	0.73	2.45	1.08	1.31	1.55	0.84		
30	PLACE1009355	5.41	7.37	9.95	13.44	10.76	13.55	*	+
	PLACE1009368	1.3	2.56	2.41	2.43	2.32	2.19		
	PLACE1009375	1.21	6.41	3.05	3.04	4.46	2.53		
	PLACE1009388	1.18	8.68	3.01	3.46	4.53	2.72		
35	PLACE1009398	1.19	9.2	3.74	3.17	4.28	3.96		
	PLACE1009404	2.78	9.18	4.51	5.33	6.73	6.94		
	PLACE1009410	1.27	2.35	2.33	2.51	2.31	1.44		
	PLACE1009417	0.95	2.25	4.34	2.55	3.08	1.71		
40	PLACE1009424	1.88	3.61	3.18	2.85	3.24	3.93		
	PLACE1009434	0.84	3.94	2.91	1.29	1.82	2.19		
	PLACE1009443	1.21	7.2	2.55	2.42	3.43	3.17		
45	PLACE1009444	1.33	7.71	4.05	2.51	3.17	3.79		
	PLACE1009459	0.23	7.99	1.55	1.71	1.83	0.86		
	PLACE1009460	1.75	6.84	3.26	5.15	4.31	4.08		
	PLACE1009468	0.99	2.83	3.42	4.43	4.42	2.97		
50	PLACE1009476	0.21	1.21	0.73	1.05	0.67	1.33		
	PLACE1009477	1.35	3.13	2.67	3.06	2.35	2.2		
	PLACE1009493	0.87	3.35	0.94	1	1.87	1.41		
	PLACE1009502	0.76	4.64	2.13	1.19	1.89	1.66		
55	PLACE1009524	1.32	4.22	1.63	0.94	2.14	1.6		

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	PLACE1008854	0.56	0.48	1.14	1.16	1.41	1.6		
	PLACE1008864	1.98	1.92	2.96	2.73	2.65	2.49		
5	PLACE1008867	1.2	6.57	6.22	6.43	5.71	4.82		
	PLACE1008876	11.7	16.5	27.29	26.74	20.94	23.24		
	PLACE1008887	1.37	4.31	1.44	3.26	2.07	3.28		
10	PLACE1008902	1.33	5.62	2.93	3.17	4.58	2.06		
	PLACE1008911	4.04	8.56	10.48	11.31	13.99	15.66		
	PLACE1008917	0.6	4.53	2.72	1.7	2.9	1.71		
	PLACE1008920	0.75	0.77	0.87	0.61	1.58	1.44		
15	PLACE1008925	0.25	0.9	1.04	0.94	1.91	0.84		
	PLACE1008930	4.12	7.32	9.83	5.11	10.36	7.17		
	PLACE1008934	0.9	3.42	2.9	2.89	2.28	1.7		
	PLACE1008941	1.57	4.14	2.8	2.06	2.59	4.05		
20	PLACE1008947	2.3	5.41	5.51	3.96	5.84	5.16		
	PLACE1008984	1.26	6.31	3.25	3.1	3.93	3.19		
	PLACE1008985	0.94	2.75	2.74	2.84	2.43	2.7		
25	PLACE1008994	0.27	1.72	0.65	1.11	0.78	0.68		
	PLACE1009020	0.46	3.49	2.42	2.49	3.1	2.16		
	PLACE1009027	0.89	2.7	1.59	2.24	1.75	2.09		
	PLACE1009039	-0.06	3.31	3.42	2.39	1.59	1.49		
30	PLACE1009045	1.53	6.33	6.05	23.13	20.76	22.2	**	+
	PLACE1009048	0.41	5.97	2.3	0.61	1.04	0.54		
	PLACE1009050	0.97	4.9	1.68	1.07	1.47	1.55		
35	PLACE1009060	5.61	8.4	9.51	10.74	8.55	11.96		
	PLACE1009067	1.14	2.8	2.03	1.6	2.34	3.4		
	PLACE1009071	1.44	4.05	3.9	3.79	7.24	9.82		
	PLACE1009090	1.27	6.46	2.35	3.11	4.73	2.86		
40	PLACE1009091	5.58	10.22	38.11	38.77	49.35	36.29		
	PLACE1009094	0.26	5.68	1.88	1.67	5.04	1.71		
	PLACE1009099	1	5.52	3.47	3.49	3.36	3.84		
	PLACE1009110	1.59	5.82	1.16	1.68	4.3	4.39		
45	PLACE1009111	1.88	5.24	2.65	3.95	3.77	2.88		
	PLACE1009113	2.24	3.52	3.62	6.14	4.87	7.29	*	+
	PLACE1009130	4.46	6.8	7.84	6.68	9.36	10.47		
50	PLACE1009150	0.88	3.54	1.95	3.23	3.3	3.01		
	PLACE1009155	1.11	5.06	2.98	4.46	4.43	3.87		
	PLACE1009158	1.06	5.77	1.95	1.77	2.35	2.88		
	PLACE1009166	0.76	4.8	1.53	1.59	2.16	1.3		
55	PLACE1009172	1.43	3.96	2.45	2.26	5.85	2.61		

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	PLACE1008533	2.01	4	4.07	5.53	5.18	3.77		
	PLACE1008542	1.61	2.36	0.96	2.05	2.1	1.72		
5	PLACE1008549	0.96	3.06	0.67	1.45	2.1	1.53		
	PLACE1008560	1.18	4.23	2.28	2.29	3.93	3.47		
	PLACE1008567	0.87	7.26	1.85	2.33	3.67	2.38		
10	PLACE1008568	2.37	10.67	5.49	2.97	7.47	4.21		
	PLACE1008569	3.94	10.32	6.74	6.1	8.6	7.6		
	PLACE1008584	0.88	1.4	1.58	2.86	3.38	1.31		
	PLACE1008585	4.96	4.8	7.56	11.08	4.84	3.57		
15	PLACE1008603	5.9	7.25	31	30.55	43.67	29.76		
	PLACE1008621	0.55	2.28	0.95	0.72	1.89	1.16		
	PLACE1008625	0.64	4.01	0.9	1.18	1.41	2.03		
20	PLACE1008626	0.55	6.06	0.9	1.03	0.83	0.95		
	PLACE1008627	0.46	8.32	1.86	1.87	3.34	2.7		
	PLACE1008629	3.22	9.18	5.84	5.44	6.75	4.41		
	PLACE1008630	1.68	3.39	4.23	4.21	3.01	3.51		
25	PLACE1008643	1.31	0.93	1.98	1.72	2.34	1.94		
	PLACE1008650	0.25	3.05	1.62	2.23	1.63	1.24		
	PLACE1008657	1.17	2.39	2.51	2.34	4.04	2.91		
	PLACE1008664	0.91	5.93	2.37	2.91	2.51	1.13		
30	PLACE1008693	0.97	4.93	3.09	2.53	3.81	2.2		
	PLACE1008696	0.88	3.84	2.21	2.26	2.11	1.47		
	PLACE1008715	1.05	4.71	2.11	1.34	2.65	2.65		
35	PLACE1008716	2.48	3.94	4.19	5.75	6.9	7.07	*	+
	PLACE1008722	3.85	4.34	7.37	7.64	7.19	3.45		
	PLACE1008738	5.17	9.13	12.7	9.49	5.83	4.8		
	PLACE1008742	6.57	6.87	14.66	14.94	15.06	12.41		
40	PLACE1008744	3.52	6.98	5.61	5.83	4.55	2.74		
	PLACE1008748	0.63	4.39	2.75	2.44	1.67	1.61		
	PLACE1008757	0.99	4.74	4.51	2.77	5.74	2.17		
	PLACE1008766	2.66	6.75	3.77	3.51	6.47	4.06		
45	PLACE1008785	1.39	1.68	2.6	3.26	3.8	3.89	*	+
	PLACE1008790	1.57	1.8	2.29	3.5	5.39	2.96		
	PLACE1008798	1.71	3.82	4.45	6	5.93	3.32		
50	PLACE1008807	1.34	3.95	1.61	2.54	2.62	1.8		
	PLACE1008808	1.6	4.53	3.01	4.24	3.69	5.04		
	PLACE1008813	1.38	4.85	1.97	1.9	1.95	2.3		
	PLACE1008836	1.34	5.81	3.68	3.89	6.17	4.1		
55	PLACE1008851	1.21	6.65	3.94	4.85	10.04	4.54		

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	PLACE1008275	1.59	3.67	2.17	2.62	2.57	2.34		
	PLACE1008280	0.85	2.6	1.84	2.42	2.48	2.36		
5	PLACE1008282	4.71	8.19	6.89	7.27	9.02	6.38		
	PLACE1008297	2.32	4.7	3.36	2.89	3.42	3.21		
	PLACE1008303	1.65	6.68	1.24	4.12	3.83	2.65		
	PLACE1008309	0.43	6.52	0.82	1.77	1.5	1.29		
10	PLACE1008315	5.3	5.93	8.61	4.92	4.79	9.83		
	PLACE1008329	0.47	2.23	2.06	2.32	2.8	2.49		
	PLACE1008330	0.72	4.06	3.16	2.48	3.36	2.96		
15	PLACE1008331	0.84	5.01	2.1	4.5	2.17	2.91		
	PLACE1008351	4.34	8.66	7.41	7.91	7.31	7.1		
	PLACE1008356	1.56	8.23	1.93	2.86	4.16	3.35		
	PLACE1008359	1.57	4.11	2.89	2	2.97	2.94		
20	PLACE1008368	2.27	6.38	7.43	5.72	7.33	6.95		
	PLACE1008369	0.57	2.46	1.45	1.12	1.59	1.68		
	PLACE1008392	0.8	3.09	2.54	2.44	3.22	3.24		
25	PLACE1008394	2.08	4.84	3.75	3.98	5.03	4.76		
	PLACE1008398	5.32	9.36	11.44	11.36	11.3	12.33		
	PLACE1008401	1.19	7.06	3.21	2.82	3.43	3.33		
	PLACE1008402	3.21	6.45	7.2	7.23	10.15	9.26		
30	PLACE1008405	10.3	10.95	18.42	17.17	18.82	20.4		
	PLACE1008409	1.88	5.19	5.69	4.97	5.41	5.65		
	PLACE1008420	1.4	1.87	1.96	2.67	2.69	2.27	*	+
35	PLACE1008424	0.88	2.69	2.54	1.69	2.34	1.71		
	PLACE1008426	0.98	2.58	1.58	1.7	2.66	2.32		
	PLACE1008429	0.92	3.17	2.14	1.91	3.4	1.84		
	PLACE1008430	1.63	4.85	3.04	2.93	3.52	3		
40	PLACE1008437	0.87	3.64	3.01	2.83	1.82	1.57		
	PLACE1008453	1.16	4.8	1.02	1.64	2.06	1.17		
	PLACE1008454	2.14	6.46	9.23	5.46	9.02	5.92		
	PLACE1008455	2.06	4.33	7.2	5.26	6.68	4.87		
45	PLACE1008457	0.51	2.6	2.01	2.28	2.43	2.47		
	PLACE1008465	0.49	2.41	1.72	1.56	2.13	0.48		
	PLACE1008469	2.42	4.36	5.32	5.16	4.75	7.1		
50	PLACE1008488	0.81	5.48	1.97	2.44	2.03	1.8		
	PLACE1008519	1.48	10.85	6.17	4.41	4.99	4.51		
	PLACE1008524	1.04	11.09	1.72	2	3.22	2.02		
	PLACE1008531	0.64	8.37	1.92	1.33	2.05	1.72		
55	PLACE1008532	2.12	3	5.51	5.66	4.72	4.19		

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	PLACE1007954	-0.03	2.45	1.15	2.46	1.57	1.51		
	PLACE1007955	0.92	4.01	2.17	2.05	2.52	3.05		
5	PLACE1007956	0.6	3.61	2.91	2.35	2.1	2.22		
	PLACE1007958	0.75	6.31	1.34	0.79	1.21	0.8		
	PLACE1007965	0.64	5.88	3.25	3.38	2.91	2.17		
10	PLACE1007969	1.09	6.37	3.06	2.35	3.29	2.21		
	PLACE1007971	2.73	4.17	5.21	6.1	4.41	5.92		
	PLACE1007990	1.95	2.33	2.31	3.22	3.09	1.88		
	PLACE1008000	0.32	2.16	1.98	1.85	1.27	0.66		
15	PLACE1008002	0.99	3.38	1.7	1.81	2.04	0.51		
	PLACE1008037	0.57	4.19	1.7	4.59	2.86	2.02		
	PLACE1008044	1.42	5.81	2.46	4.18	4.93	4.16		
20	PLACE1008045	0.4	4.07	1.54	1.75	2	1.65		
	PLACE1008080	2.05	6.08	3.22	4.23	4.03	4.78		
	PLACE1008092	1.56	1.56	1.48	1.48	2.98	1.86		
	PLACE1008095	0.59	2.14	1.48	2.38	2.73	1.23		
25	PLACE1008105	0.95	1.76	1.71	2.24	2.71	0.74		
	PLACE1008107	0.27	2.33	0.7	1.72	1.44	1.68		
	PLACE1008111	1.73	5.01	2.12	4.57	5.4	4.04		
30	PLACE1008113	5.88	9.24	12.48	16.57	20.29	19.24	*	+
	PLACE1008122	1.22	5.54	2.55	1.61	1.57	1.5		
	PLACE1008129	1.5	5.64	2.8	2.43	5.36	2.91		
	PLACE1008132	5.51	4.47	8.34	6.61	11.2	7.63		
35	PLACE1008137	0.96	1.82	1.02	2.12	3.88	0.8		
	PLACE1008174	0.77	3.16	2.43	5.12	4.39	2.46		
	PLACE1008177	1.62	4.87	3.09	3.79	3.26	3.77		
	PLACE1008181	1.76	3.87	1.6	2.06	2.43	1.43		
40	PLACE1008195	2.66	6.08	2.97	4.34	4.14	3.9		
	PLACE1008198	1.06	5.56	2.52	2.55	3.31	1.54		
	PLACE1008201	1.22	4.45	3.58	5.92	7.69	5.64	*	+
45	PLACE1008209	2.35	2.29	4.46	2.51	5.31	4.44		
	PLACE1008226	1.8	2.35	5.25	4.72	5.68	7.08		
	PLACE1008227	0.77	2.67	3.02	2.38	4.6	3.54		
	PLACE1008231	1.26	3.85	1.85	1.05	1.83	0.78		
50	PLACE1008238	1.22	3.21	2.9	2.47	2	1.72		
	PLACE1008244	1.01	4.69	1.56	1.68	3.2	1.56		
	PLACE1008249	0.8	4.94	2.55	1.22	2.17	2.01		
	PLACE1008266	11.31	18.61	43.04	60.04	82.48	59.04	*	+
55	PLACE1008273	1.47	3.95	3.81	3.53	3.8	4.47		

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	PLACE1007690	1.4	4.03	2.12	3.74	3.37	4.61		
	PLACE1007697	0.69	7.13	1.37	1.84	2.56	1.8		
5	PLACE1007702	2.03	7.08	5.7	4.03	3.91	4.08		
	PLACE1007705	1.38	3.93	1.59	1.74	4.45	2.75		
	PLACE1007706	3.11	6.08	4.69	5.25	8.84	7.49		
	PLACE1007725	3.41	4.69	6.65	5.1	3.29	5.39		
10	PLACE1007729	0.98	2.65	1.8	2.7	3.11	1.99		
	PLACE1007730	1.25	4.29	3.07	3.66	3.75	4.32		
	PLACE1007737	1.43	4.79	3.39	3.79	4.7	4.17		
15	PLACE1007743	1.38	4.26	2.29	2.3	2.83	2.03		
	PLACE1007746	6.56	9.02	10.42	9.65	13.29	12.97		
	PLACE1007753	0.53	4.48	1.71	1.35	2.86	1.94		
	PLACE1007769	1.31	4.31	3.5	3.27	4.51	4.58		
20	PLACE1007780	5.77	4.63	7.11	6.51	3.75	2.17		
	PLACE1007791	1.82	3.29	3.38	3.16	3.69	2.87		
	PLACE1007807	0.67	2.79	1.72	2.33	1.76	1.29		
25	PLACE1007810	0.39	4.45	2.63	4.11	4.08	3.27		
	PLACE1007814	3.57	5.98	5.04	4.2	4.62	6.3		
	PLACE1007828	2.01	7.64	3.34	2.69	4.64	3.44		
	PLACE1007829	1.32	6.9	2.88	2.87	4.87	3.06		
30	PLACE1007841	1.64	7.26	1.87	2.25	3.14	3.39		
	PLACE1007842	1.1	3.32	2.44	2.09	3.96	1.39		
	PLACE1007843	1.2	1.92	1.43	2.13	1.48	1.86		
35	PLACE1007845	1.76	3	4.11	3.45	3.42	2.36		
	PLACE1007846	0.99	3.26	1.64	2.02	2.73	1.5		
	PLACE1007848	1.09	3.51	2.23	2.39	2.62	2.25		
	PLACE1007852	2.26	7.88	3.82	2.94	4.61	3.24		
40	PLACE1007858	3.65	11.57	5.81	61.71	80.46	57.09	**	+
	PLACE1007866	19.42	25.98	40.48	43	80.39	56.73		
	PLACE1007871	8.1	7.9	15.45	16.17	12.35	11.08		
	PLACE1007877	1.09	2.09	1.45	1.4	2.39	1.53		
45	PLACE1007878	5.98	9.75	14.61	13.65	7.49	8.9		
	PLACE1007881	0.43	2.66	1.34	1.59	1.94	1.93		
	PLACE1007885	4.35	7.85	6.76	5.57	6.53	7.01		
50	PLACE1007897	0.27	6.51	1.85	1.72	1.53	1.41		
	PLACE1007908	3.14	12.29	5.73	5.96	7.9	8.24		
	PLACE1007922	6.08	11.75	8.75	5.24	7.15	4.54		
	PLACE1007946	1.07	2.03	1.86	2.71	2.28	1.94		
55	PLACE1007950	6.98	7.6	18.21	16.17	19.34	12.63		

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	PLACE1007402	2.41	5.66	3.08	1.52	2.91	1.8		
	PLACE1007409	1.05	4.57	1.04	2.51	2.68	2.02		
5	PLACE1007416	3.45	6.97	6.5	7.05	9.14	6.52		
	PLACE1007420	12.12	12.66	20.8	25.26	23.9	22.88	*	+
	PLACE1007431	1.87	7.4	5.51	7.17	5.28	5.91		
	PLACE1007450	0.79	1.22	2.65	3	2.99	2.39		
10	PLACE1007452	0.42	2.36	1.76	2.09	2.98	1.45		
	PLACE1007454	23.74	28.02	76.56	59.97	75.95	46.61		
	PLACE1007460	0.75	3.52	2.35	2.34	1.93	2.58		
15	PLACE1007478	0.41	3.07	1.33	1.35	2.18	1.92		
	PLACE1007484	0.6	4.8	2.57	2.56	1.45	1.69		
	PLACE1007488	0.4	6.24	1.64	1.74	2.61	1.46		
	PLACE1007507	2.91	6.36	4.49	5.31	8.29	8.11		
20	PLACE1007511	0.53	1.29	1.06	1.06	1.29	0.42		
	PLACE1007513	10.57	10.43	24.05	12.24	16.88	16.9		
	PLACE1007524	1.55	3.33	3.53	3.96	4.72	2.96		
25	PLACE1007525	1.24	2.95	3.14	2.38	2.85	2.24		
	PLACE1007537	8.6	9.68	49.88	43.78	63.66	40.1		
	PLACE1007544	1.55	6.45	4.97	3.2	3.92	4.61		
	PLACE1007547	1.36	5.03	4.15	2.37	2.84	2.4		
30	PLACE1007557	1.12	3.16	3.14	3.07	3.9	3.41		
	PLACE1007560	9.38	8.86	12.57	11.03	9.62	17.59		
	PLACE1007565	0.37	2.27	1	1	1.16	0.91		
35	PLACE1007580	1.06	3.71	3.06	10.8	11.15	13.74	**	+
	PLACE1007583	0.76	3.88	1.78	2.51	2.37	1.09		
	PLACE1007591	0.79	4.62	1.7	2.2	2.53	2.07		
	PLACE1007598	1.13	6.98	3.86	2.71	3.46	4.71		
40	PLACE1007610	0.41	5.63	1.28	1.33	3.18	1.5		
	PLACE1007618	1.57	1.91	2.01	1.75	2.2	2.41		
	PLACE1007621	1.78	2.83	3.64	3.33	3.57	4.38		
	PLACE1007626	23.99	25.61	32.78	30.53	30.94	13.53		
45	PLACE1007632	2.03	3.26	2.52	2.65	3.81	4.63		
	PLACE1007635	1.61	4.62	6.42	2.8	4.19	3.37		
	PLACE1007645	10.59	11.55	15.06	9.99	11.58	11.95		
50	PLACE1007649	1.7	5.88	3.47	2.78	4.95	3.13		
	PLACE1007659	1.33	5.85	3.61	4.88	6.22	4.9		
	PLACE1007669	2.01	2.1	3.74	2.97	4.63	4.4		
	PLACE1007677	1.25	2.29	2.81	2.68	3.07	2.91		
55	PLACE1007688	3.4	5.69	5.43	1.98	4.53	4.98		

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	PLACE1007105	1.18	6.51	3.44	2.65	4.13	2.21		
	PLACE1007108	3.55	13.02	7.41	5.03	6.87	5.75		
5	PLACE1007111	1.33	9.51	-1.52	1.74	2.37	1.52		
	PLACE1007112	1.23	7.26	1.79	2.09	3.12	2.36		
	PLACE1007130	0.54	2.02	1.92	0.87	1.47	0.33		
	PLACE1007132	1.46	3.32	4.63	3.58	3.38	2.88		
10	PLACE1007140	0.61	2.58	2.41	1.98	1.98	1.32		
	PLACE1007143	2.79	6.32	4.62	4.9	5.34	5.33		
	PLACE1007169	2.21	8.59	3.46	5.44	8.46	7.99		
15	PLACE1007178	0.82	8.66	2.48	3.28	6.28	4.1		
	PLACE1007190	3.31	10.9	6.7	10.51	13.57	11.14		
	PLACE1007201	0.81	5.82	1.41	1.72	3.04	2.51		
	PLACE1007202	37.76	34.95	76.28	58.23	34.42	37.86		
20	PLACE1007226	2.01	2.39	2.73	1.89	3.14	2.29		
	PLACE1007238	1.64	3.07	1.83	2.39	2.73	2.2		
	PLACE1007239	1.81	3.68	2.99	1.76	2.72	2.44		
25	PLACE1007242	0.61	5.18	1.87	1.54	1.14	1.67		
	PLACE1007243	2.21	7.36	2.29	2.24	3.27	3.31		
	PLACE1007247	0.36	6.17	1.71	1.11	1.36	1.34		
	PLACE1007257	1.67	5.33	3.34	3.3	5.27	4.25		
30	PLACE1007274	1.46	2.18	4.43	4.38	4.03	4.06		
	PLACE1007276	0.93	2.02	1.1	2.13	2.1	1.74		
	PLACE1007282	2.51	4.2	5.72	4.28	3.62	4.66		
35	PLACE1007286	2.97	4.8	7.85	10.14	12.47	8.79	*	+
	PLACE1007296	10.55	19.45	24.46	31.43	17.57	27.05		
	PLACE1007301	0.65	5.17	1.55	1.19	1.54	1.11		
	PLACE1007314	3.11	6.61	8.64	7.98	8.96	10.24		
40	PLACE1007317	1.19	3.34	1.27	1.88	1.62	1.79		
	PLACE1007329	0.89	0.73	1.78	2.38	2.35	2.09	*	+
	PLACE1007338	3.96	6.47	9.58	11.59	8.93	2.32		
	PLACE1007342	0.71	1.8	1.3	1.1	1.1	0.7		
45	PLACE1007345	1.72	4.57	2.54	2.72	3.6	3.29		
	PLACE1007346	1.43	4.61	3.89	5.77	4.53	4.1		
	PLACE1007359	0.74	4.55	2.16	2.59	2.44	3.53		
50	PLACE1007367	4.53	8.63	15.16	12.49	13.49	11.75		
	PLACE1007375	0.36	3.24	2.02	1.75	2.56	1.59		
	PLACE1007377	1.49	2.01	3.18	3.29	3.96	2.36		
	PLACE1007386	1.55	1.75	1.47	2.37	1.68	1.36		
55	PLACE1007392	1.57	2.99	2.49	2.79	4.48	3.51		

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	PLACE1006860	1	4.29	1.62	1.61	2.1	1
	PLACE1006867	5.65	9.36	11.34	7.04	8.33	7.63
5	PLACE1006875	1.15	6.19	5.66	4.84	4.53	4.63
	PLACE1006878	1.59	2.84	3.09	2.99	3.22	2.39
	PLACE1006883	3.21	5.08	6.78	6.83	7.38	6.19
	PLACE1006898	1.67	4.23	3.67	3.54	4.77	4.59
10	PLACE1006901	2.59	4.75	4.03	3.71	3.28	4.14
	PLACE1006904	0.91	3.59	2.7	3.26	2.92	2.04
	PLACE1006917	3.63	7.13	6.1	5.8	7.21	7.03
15	PLACE1006932	0.54	5.85	1.29	0.92	1.34	1.19
	PLACE1006935	1.3	5.46	2.54	1.59	4.03	1.6
	PLACE1006956	0.92	2.55	3.4	2.55	2.41	2.09
	PLACE1006958	0.78	2.41	1.35	1.76	4.2	3.39
20	PLACE1006959	4.97	8.48	9.98	11.46	9.58	13.62
	PLACE1006961	8.03	9.85	14.42	13.73	11.57	14.2
	PLACE1006962	2.97	7.44	6.56	5.04	7.26	6.22
25	PLACE1006966	2.02	6.94	3.46	3.15	3.89	2.89
	PLACE1006979	0.95	4.44	2.03	1.46	2.64	1.77
	PLACE1006989	2.19	5.05	3.02	3.27	3.9	5.06
	PLACE1007001	4.98	6.79	10.71	4.03	7.43	7.38
30	PLACE1007014	1.37	3.03	3.45	1.79	2.18	2.2
	PLACE1007021	0.74	3.03	2.11	0.75	2.2	1.73
	PLACE1007026	2.1	9.23	3.93	4.15	4.27	5.42
	PLACE1007028	4.12	8.5	10.56	7.89	8.34	9.35
35	PLACE1007038	237.33	267.91	446.14	406.27	622.67	671.17
	PLACE1007040	1.55	3.14	2.85	1.57	3.31	2.45
	PLACE1007045	1.08	3.74	2.85	2.9	5.03	2.74
40	PLACE1007048	147.06	149.67	259.53	121.61	211.26	109.43
	PLACE1007053	4.9	6.69	10	3.59	4.91	4.71
	PLACE1007068	7.56	10.33	62.76	39.52	45.9	36.69
	PLACE1007070	5.97	10.85	10.28	8.65	9.6	14.3
45	PLACE1007076	8.22	14.4	14.19	16.53	23.62	24.67 * +
	PLACE1007077	2.65	6.45	4.01	5.2	5.28	5.43
	PLACE1007081	0.36	4.47	1.94	1.92	1.92	1.37
50	PLACE1007082	1.23	4.66	4.95	4.32	4.5	3.99
	PLACE1007092	2.49	4.12	7.26	4.77	5.22	4.34
	PLACE1007096	0.72	2.19	0.74	1.35	1.63	0.97
	PLACE1007097	0.54	2.49	1.35	1.61	1.28	1.04
55	PLACE1007099	1.58	4.66	2.56	2.77	3.64	3.72

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	PLACE1006607	1.47	7.69	4.18	3.45	5.86	4.29		
	PLACE1006610	9.46	13.73	38.26	27.65	32.76	22.64		
5	PLACE1006615	6.22	9.09	48.78	20.25	15.74	15.86		
	PLACE1006617	0.91	1.54	2.66	1.87	2.49	2.09		
	PLACE1006618	5.42	8.01	9.24	5.33	8.59	5.76		
	PLACE1006626	1.53	4.11	1.3	2.47	2.78	1.16		
10	PLACE1006629	0.99	5.05	1.36	2.22	2.56	1.76		
	PLACE1006637	1.29	6.54	3.97	3.77	4.23	4.87		
	PLACE1006640	0.59	5.14	1.17	0.85	2.54	0.94		
15	PLACE1006644	1.66	4.46	2.12	2.79	2.49	2.39		
	PLACE1006657	1.28	2.09	2.31	4.55	3.09	2.19		
	PLACE1006673	2.29	4.73	10.34	11.06	10.89	6.45		
	PLACE1006678	2.54	2.98	1.44	1.37	1.96	1.39		
20	PLACE1006682	3.5	5.93	2.58	15.44	20.96	23.99	**	+
	PLACE1006684	1.12	4.8	1.81	1.64	2.54	1.65		
	PLACE1006698	1.54	5.86	4.52	2.15	3.57	1.9		
25	PLACE1006704	1.81	5.41	2.71	2.93	2.92	2.97		
	PLACE1006708	1.69	5.07	3.49	3.46	4.11	3.7		
	PLACE1006711	14.21	16.18	29.77	24.34	26.25	22.42		
	PLACE1006714	2.27	3.26	4.74	4.57	5.23	3.53		
30	PLACE1006716	1.51	2.75	3.7	6	7.05	3.99		
	PLACE1006731	1.65	3.77	2.83	2.71	4	3.09		
	PLACE1006754	0.43	3.94	1.73	1.8	1.81	0.99		
35	PLACE1006760	7.56	10.98	10.08	8.58	8.89	11.31		
	PLACE1006779	1.44	4.12	2.88	3.19	3.79	2.97		
	PLACE1006782	0.44	5.17	2.42	2.95	1.57	1.15		
	PLACE1006783	9.34	11.46	18.65	157.98	223.05	66.46	*	+
40	PLACE1006786	3.31	4.08	6.07	5.9	6.24	3.34		
	PLACE1006792	1.61	3.31	5.38	5.66	3.33	4.18		
	PLACE1006795	0.89	2.43	0.74	0.81	1.27	1.01		
	PLACE1006800	1.62	4.94	2.53	4.7	4.56	3.93		
45	PLACE1006805	3.94	7.79	5.5	10.83	9.8	8.79	*	+
	PLACE1006809	3.55	5.7	5.94	9.58	10.61	8.97	**	+
	PLACE1006815	1.7	7.57	4.1	5.12	5.23	5.8		
50	PLACE1006819	0.33	0.88	0.95	0.89	1.76	0.63		
	PLACE1006820	2.35	2.01	4.91	4.84	6.72	4.18		
	PLACE1006826	2.28	6.22	4.84	7.68	7.62	5.58		
	PLACE1006829	3.76	5.51	6.54	9.49	8.66	8.69	*	+
55	PLACE1006853	1.2	4.21	1.97	2.25	2.93	2.88		

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	PLACE1006365	1.68	4.65	1.97	1.8	2.12	0.97		
	PLACE1006368	1.53	3.11	2.57	3.01	3.04	4.27		
5	PLACE1006371	1.38	3.2	1.46	1.68	3.01	1.67		
	PLACE1006373	2.21	5.21	5.75	7.83	8.02	7.56	*	+
	PLACE1006382	0.9	4.67	2.81	3.3	1.92	2.95		
	PLACE1006385	1.59	6.33	1.86	2.68	2.59	2.71		
10	PLACE1006391	1.19	5	1.95	1.96	2.79	1.63		
	PLACE1006412	1.88	5.53	5.92	7.07	9.93	5.27		
	PLACE1006414	0.63	3.42	0.95	1.22	1.87	1.6		
15	PLACE1006419	7.79	9.8	11.93	5.19	7.29	5.32	*	-
	PLACE1006438	0.99	6.07	3.42	3.29	4.56	5.14		
	PLACE1006443	2.05	5.01	5.12	5.01	5.31	6.44		
	PLACE1006445	0.84	5.76	3.65	3.53	3.27	3.55		
20	PLACE1006447	1.34	5.81	3.28	2.95	3.26	3.96		
	PLACE1006466	0.75	4.38	1.35	1.49	1.66	1.08		
	PLACE1006469	0.67	4.66	2.31	1.65	2.26	1.67		
25	PLACE1006470	2.47	3.71	3.74	5.25	7.02	4.35		
	PLACE1006472	24.4	23.44	52.17	26.23	28.52	9.36		
	PLACE1006476	2.52	4.31	8.67	6.21	7.23	5.93		
	PLACE1006482	1.64	3.35	4.43	4.25	4.67	4.98		
30	PLACE1006488	14.12	19.42	32.69	40.76	34.77	41.4	*	+
	PLACE1006492	2.03	6.41	4.38	4.04	4.98	3.02		
	PLACE1006506	1.78	6.67	4.04	4.41	5.71	4.17		
35	PLACE1006515	1.65	5.7	3.08	3.19	2.84	4.08		
	PLACE1006516	1.1	7.32	7.05	4.89	5.69	7.28		
	PLACE1006520	1.02	2.74	2.12	1.19	3	1.54		
	PLACE1006521	2.4	3.54	6.38	6.49	6.86	5.08		
40	PLACE1006529	5.96	7.35	6.96	10.56	8.2	7.93		
	PLACE1006531	1.01	4.31	3.33	1.84	2.05	2.43		
	PLACE1006534	1.68	6.04	2.59	3.01	3.86	3.19		
45	PLACE1006540	2.68	9.7	7.77	8.71	11.21	4.46		
	PLACE1006549	0.6	9.45	2.09	1.6	2.28	1.65		
	PLACE1006550	1.76	8.82	4.07	2.77	2.94	4.14		
	PLACE1006552	1.3	2.48	2.14	1.97	1.3	0.81		
50	PLACE1006557	2.38	4.01	3.79	2.84	2.51	2.45		
	PLACE1006563	2.49	3.44	5.7	4.23	4.15	4.3		
	PLACE1006579	1.53	7.5	4.82	4.88	5.38	5.78		
	PLACE1006594	236.53	241.11	397.64	122.37	278.58	324.29		
55	PLACE1006598	0.72	8.53	2.4	1.53	1.58	2.07		

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	PLACE1006139	3.44	2.98	6.03	7.77	8.85	5.58		
	PLACE1006143	0.5	1.48	1.87	3.18	4.13	3.17	*	+
5	PLACE1006157	1.55	2.54	-4.82	2.96	3.9	2.44		
	PLACE1006159	0.69	3.61	0.94	2.68	1.98	1.04		
	PLACE1006164	0.35	3.18	1.37	1.73	1.85	1.21		
	PLACE1006167	2.18	6.5	3.37	3.95	4.52	3.13		
10	PLACE1006170	2.79	6.09	6.09	4.34	5.31	3.68		
	PLACE1006181	2.75	7.34	2.84	5.8	5.51	5.22		
	PLACE1006187	0.76	1.3	2.15	2.01	2.41	1.48		
15	PLACE1006195	0.11	1.24	1.73	1.93	1.93	0.87		
	PLACE1006196	1.8	4.01	4.15	4.32	5.77	2.19		
	PLACE1006197	2.12	5.6	5.24	4	3.47	3.39		
	PLACE1006198	0.27	3.68	1.21	0.84	1.63	0.5		
20	PLACE1006205	0.89	5.59	0.99	2.43	2.18	1.28		
	PLACE1006208	7.28	13.32	13.46	14.09	14.99	12.5		
	PLACE1006211	2.6	8.05	7.92	6.07	9.08	9.35		
25	PLACE1006219	6.77	5.77	8.94	14.88	22.25	15.35	*	+
	PLACE1006223	1.55	1.46	3.19	1.39	3	1.64		
	PLACE1006225	0.56	2.27	1.3	1.04	2.3	0.99		
	PLACE1006236	1.53	3.2	2.92	3.06	5.01	2.29		
30	PLACE1006239	0.67	3.62	1.97	2.61	3.66	3.41		
	PLACE1006245	3.86	7.13	5.45	4.43	7.44	3.28		
	PLACE1006246	1.66	6.56	6.19	5.59	7.66	6.33		
	PLACE1006248	1.58	4.47	5.6	2.77	3.1	2.82		
35	PLACE1006262	0.93	2.24	1.49	2.08	1.61	1.4		
	PLACE1006269	2.28	4.71	3.42	2.06	2.47	2.33		
	PLACE1006275	1.6	3.57	3.37	4.12	3.68	3.53		
40	PLACE1006277	1.01	2.42	1.4	1.79	3.01	0.88		
	PLACE1006288	9.32	13.59	22.49	26.85	18.4	25.21		
	PLACE1006290	1.79	6.81	5.99	8.87	7.56	9.13		
	PLACE1006298	1.93	5.52	2.47	3.87	5.08	4.55		
45	PLACE1006311	0.65	3.38	1.75	225.97	161.43	251.12	**	+
	PLACE1006318	3.52	4.03	4.17	4.04	3.17	4.01		
	PLACE1006325	5.43	6.73	6.31	8.09	8.38	8.08	**	+
50	PLACE1006331	1.87	3.36	3.21	4.44	3.59	2.56		
	PLACE1006335	1.76	3.64	2.55	4.45	2.98	2.92		
	PLACE1006357	0.27	4.51	1.59	1.7	1.49	1.2		
	PLACE1006360	1.1	5.11	1.79	2.46	2.74	2.62		
55	PLACE1006364	4.51	8.06	7.29	7.37	9.19	5.75		

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	PLACE1005876	1.33	3.99	4.76	6.87	6.34	6.9	*	+
	PLACE1005878	1.3	2.67	2.08	3.54	4.46	2.79		
5	PLACE1005880	2.36	4.09	4.31	4.16	3.07	3.45		
	PLACE1005884	1.6	4.87	1.89	2.48	2.21	2.73		
	PLACE1005890	1.9	9.57	3.7	2.26	3.09	1.88		
10	PLACE1005898	3.29	10.87	5.34	7.36	8.12	6.36		
	PLACE1005913	1.46	9.31	8.05	4.99	6.47	4.33		
	PLACE1005921	0.99	1.92	2.03	1.49	2.22	0.79		
	PLACE1005923	0.74	1.61	1.17	1.47	2.42	1.43		
15	PLACE1005925	0.83	2.67	3.18	2.19	2.14	1.68		
	PLACE1005927	1.26	2.49	1.93	1.95	2.56	2.3		
	PLACE1005932	2.04	5.66	2.44	2.53	2.32	2.52		
	PLACE1005934	0.88	7.91	3.16	3.9	5.61	4.19		
20	PLACE1005936	1.31	8.96	3.02	2.02	2.84	2.3		
	PLACE1005939	54.61	68.58	111.22	157.61	194.58	212.18	**	+
	PLACE1005951	2.36	3.39	4.98	5.56	4.48	2.35		
25	PLACE1005953	1.5	1.64	2.64	2.59	2.43	3.03		
	PLACE1005955	1.64	2.01	3.8	4.07	3.43	2.55		
	PLACE1005966	0.76	3.42	1.69	1.75	2	2.19		
	PLACE1005968	1.52	4.96	3.2	4.71	5.15	6.12		
30	PLACE1005975	2.58	7.11	5.42	6.18	7.01	6.49		
	PLACE1005990	0.7	7.7	1.54	2.1	1.87	0.88		
	PLACE1005997	88.15	118.52	196.48	189.6	226.97	172.1		
35	PLACE1006002	3.38	3.97	8.87	8.4	7.71	9.18		
	PLACE1006003	1.55	3.02	4.83	5.09	4.44	5.33		
	PLACE1006011	1.85	3.63	3.46	4.48	2.68	1.91		
	PLACE1006017	0.84	2.74	2.81	3.4	3.4	3.58		
40	PLACE1006037	2.99	7.05	2.48	6.14	3.64	4.29		
	PLACE1006040	2.2	7.87	3.97	6.64	6.9	7.77		
	PLACE1006063	0.94	4.64	2.59	2.11	3.15	2.25		
45	PLACE1006071	3.06	6.52	4.97	5.36	4.03	4.47		
	PLACE1006073	2.74	3.53	6.43	7.19	6.81	6.93		
	PLACE1006074	1.4	2.22	3.34	2.62	3.23	1.69		
	PLACE1006076	1.36	2.51	2.98	3.15	2.47	2.75		
50	PLACE1006079	1.38	4.32	1.78	2.1	1.1	1.11		
	PLACE1006093	0.49	3.76	1	3.56	3.85	1.83		
	PLACE1006116	2.99	6.44	4.04	5.28	5.01	4.91		
	PLACE1006119	3.15	6.81	7.07	9.22	10.4	8.03		
55	PLACE1006129	2.12	5.6	3.98	6.59	7.62	5.65		

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	PLACE1005656	0.92	2.48	2.24	1.68	2.78	1.58		
	PLACE1005659	0.87	2.64	1.01	1.62	1.84	1.32		
5	PLACE1005660	3.91	8.03	-5.77	8.87	8.88	8.34		
	PLACE1005664	2.69	6.57	6.14	3.39	4.27	3.19		
	PLACE1005666	0.89	5.91	3.55	4.63	4.93	3.97		
	PLACE1005669	4.46	10.41	11.39	11.64	13.9	14.6		
10	PLACE1005682	1.94	5.27	4.49	6.2	5.2	5.47		
	PLACE1005698	0.6	2.7	2.92	2.01	2.67	2.38		
	PLACE1005708	25.32	34.08	53.46	53.89	59.98	53.76		
15	PLACE1005725	3.25	3.75	6.41	5.64	5.82	7.29		
	PLACE1005727	2.97	4.54	4.15	3.9	3.49	4		
	PLACE1005730	0.77	4.29	3.26	1.1	1.54	1.28		
	PLACE1005736	5.37	7.55	5.73	9.25	12.55	10.19	*	+
20	PLACE1005739	0.81	4.96	1.38	2.46	3.17	1.74		
	PLACE1005745	8.03	7.11	11.52	11.98	6.97	11.44		
	PLACE1005752	1.31	3.15	2.96	2.55	2.24	1.25		
25	PLACE1005755	0.8	2.79	3.02	1.72	3.28	2.27		
	PLACE1005756	10.79	12.06	17.2	18.22	19.3	21.47	*	+
	PLACE1005760	10.22	15.24	68.06	49.69	68.81	53.09		
	PLACE1005763	1.47	7.04	3.58	3.79	4.63	3.02		
30	PLACE1005768	1.25	5.63	3.69	4.58	5.13	4.19		
	PLACE1005771	5.71	13.63	13.7	11.28	17.49	17.27		
	PLACE1005783	1.82	2.44	3.64	3.05	3.71	3.47		
	PLACE1005799	4.79	5.25	8.37	6.12	8.78	8.62		
35	PLACE1005802	1.07	3.78	3.64	2.7	3.64	1.96		
	PLACE1005803	3.06	6.15	4.78	5.6	4.94	7.36		
	PLACE1005804	0.92	8.41	1.33	2	1.91	2.44		
40	PLACE1005813	17.23	18.71	78.06	70.01	94.17	74.89		
	PLACE1005815	1.43	5.6	4.38	3.8	5.01	4.1		
	PLACE1005828	2.11	3.62	4.42	5.34	6.24	3.56		
	PLACE1005833	119.17	92.82	182.22	122	114.37	107.96		
45	PLACE1005834	2.04	4.33	3.95	3.55	3.56	2.56		
	PLACE1005835	22.7	19.1	51.52	72.32	60.34	68.56	*	+
	PLACE1005836	2.39	4.21	4.97	2.61	3.83	2.55		
50	PLACE1005845	0.97	5.42	2.66	2.67	3.05	3.65		
	PLACE1005850	1.82	3.91	3.04	2.84	2.85	2.15		
	PLACE1005851	1.03	3.44	1.46	1.2	2.01	1.11		
	PLACE1005856	0.92	4.01	2.42	2.24	3.37	3.37		
55	PLACE1005875	1.78	3.89	4.77	3.3	3.48	3.17		

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	PLACE1005477	1.58	2.26	2.51	3	2.93	2.74		
	PLACE1005480	0.77	2.01	1.86	1.93	1.4	0.53		
5	PLACE1005481	0.44	4.81	2.3	2.77	3.62	2.44		
	PLACE1005494	0.27	6.66	1.68	1.21	1.73	1.06		
	PLACE1005495	3.86	12.83	8.31	6.85	9.25	7.62		
	PLACE1005497	2.27	7.72	3.95	4.24	5.68	5.91		
10	PLACE1005499	5.71	5.86	11.07	10.82	7.9	6.49		
	PLACE1005502	1.59	2.87	3.43	4.07	3.45	1.49		
	PLACE1005513	1.77	4.14	3.35	1.86	2.85	1.98		
15	PLACE1005515	2.89	4.76	4.22	4.58	5.29	3.78		
	PLACE1005519	1.04	4.53	3.29	2.85	2.85	2.83		
	PLACE1005526	0.58	5.55	1.38	1.3	1.59	0.71		
	PLACE1005528	2.08	7.71	5.57	5.94	7.12	5.33		
20	PLACE1005530	2.16	7.09	4.32	5.17	8.23	4.67		
	PLACE1005536	1.74	1	2.74	3.12	2.43	2.88		
	PLACE1005539	10.1	11.64	23.77	8.65	8.66	5.22		
25	PLACE1005543	1.7	3.57	5.62	3.54	4.32	2.57		
	PLACE1005544	0.86	3.26	3.15	2.49	2.68	2.27		
	PLACE1005550	4.32	7.61	7.85	10.16	7.25	6.86		
	PLACE1005554	1.15	5.47	2.67	2.17	2.17	1.17		
30	PLACE1005557	1.76	7.21	4.95	8.22	7.64	7.7		
	PLACE1005563	0.51	4	1.89	1.45	2.07	1.06		
	PLACE1005569	0.6	0.5	1.56	1.59	1.81	1.09		
35	PLACE1005574	1.07	1.88	2.49	2.48	4.43	2.22		
	PLACE1005584	1.3	2.68	3.91	3.91	5.58	3.03		
	PLACE1005590	4.28	5.14	8.4	9.87	10.73	8.02		
	PLACE1005595	3.08	4.03	2.89	3.65	3.81	3.89		
40	PLACE1005601	2	5.66	4.22	3.77	4	4.02		
	PLACE1005603	1.08	4.9	1.04	2.49	0.95	1.94		
	PLACE1005604	1.2	6.71	2.42	3.6	4.2	3.46		
	PLACE1005611	2.22	2.3	3.98	5.15	5.65	2.89		
45	PLACE1005622	0.65	1.71	1.98	2.94	3.88	1.26		
	PLACE1005623	1.42	3.08	3.27	3.71	3.65	1.61		
	PLACE1005630	3.31	5.81	7.75	87.83	72.15	89.12	**	+
50	PLACE1005639	0.75	4.36	1.28	1.66	2.02	1.18		
	PLACE1005646	2.13	5.41	4.31	5.4	5.08	2.55		
	PLACE1005647	2.77	9.69	6.72	7.34	9.11	6.25		
	PLACE1005648	3	8.11	9.21	8.34	10.59	8.22		
55	PLACE1005653	1.99	1.43	2.74	2.74	2.13	2.67		

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	PLACE1005193	1.48	3.78	1.71	3.84	2.91	2.61		
	PLACE1005200	1.35	4.68	2.61	2.47	3.75	3.1		
5	PLACE1005206	2.43	6.48	4.26	3.35	3.95	2.95		
	PLACE1005216	1.53	5.46	4.44	5.6	6.51	4.12		
	PLACE1005223	1.43	6.21	5	4.38	5.66	3.27		
	PLACE1005225	1.36	3.01	3.49	3.33	3.32	4.65		
10	PLACE1005232	1.86	3.31	4.87	5.63	6.19	3.88		
	PLACE1005239	1.06	4.3	2.32	2.84	2.86	2.41		
	PLACE1005243	4.35	7.32	5.41	8.48	7.49	10.75		
15	PLACE1005250	4.24	10.31	7.98	4.38	5.9	8.88		
	PLACE1005261	3.21	7.43	4.74	4.78	5.82	3.51		
	PLACE1005266	1.05	4.47	2.82	2.28	4.43	2.76		
	PLACE1005271	4.66	5.31	8.79	5.87	11.16	7.95		
20	PLACE1005277	2.06	3.48	2.35	2.62	1.98	2.64		
	PLACE1005287	3.63	4.31	5.87	2.98	5.06	6.91		
	PLACE1005299	24.16	22.75	48.29	35.17	24.24	41.06		
25	PLACE1005305	6.81	8.46	11.13	10.67	11.85	16.25		
	PLACE1005307	1.59	5.44	4.14	3.15	5.42	4.84		
	PLACE1005308	2.41	4.96	3.95	5.32	5.99	5.79		
	PLACE1005313	1.08	3.83	1.6	1.8	2.05	1.8		
30	PLACE1005320	1.36	3.65	3.34	3.39	4.05	2.26		
	PLACE1005327	10.78	8.74	16.8	10.36	7.95	4.43		
	PLACE1005331	2.28	4.92	5.28	4.66	4.97	3.33		
	PLACE1005335	1.53	3.8	2.24	2.03	3.22	2.42		
35	PLACE1005336	9.12	12.58	16.58	16.39	16.99	20.15		
	PLACE1005351	2.62	8.18	10.17	9.28	8.66	9.52		
	PLACE1005366	2.04	6.93	3	2.99	3.71	4.23		
40	PLACE1005373	1.77	6.34	4.44	3.91	5.36	3.37		
	PLACE1005374	3.29	9.47	11.4	7.35	10.22	12.41		
	PLACE1005383	8.16	7.54	12.81	7.21	5.93	4.03		
	PLACE1005388	0.33	2.04	1.56	1.92	3.67	2.2		
45	PLACE1005409	2.97	5.02	4.99	3.9	4.23	2.97		
	PLACE1005410	12.41	16.44	18.89	24.38	20.98	27.1	*	+
	PLACE1005426	5.16	7.48	9.06	5.51	7.67	5.45		
50	PLACE1005431	12.6	15.65	22.53	19.64	26.25	23.75		
	PLACE1005453	1.4	10.38	3.93	4.85	4.45	3.28		
	PLACE1005467	3.09	11.87	7	5.57	11.63	7.28		
	PLACE1005471	1.6	1.94	1.66	2.29	1.52	1.28		
55	PLACE1005476	0.42	1.73	1.24	1.6	1.57	1.46		

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	PLACE1005004	0.47	3.36	0.92	1.5	0.73	0.85		
	PLACE1005005	3.35	6.9	5.32	6.67	9.65	7.17		
5	PLACE1005011	6.03	11.12	-35.8	37.66	56.97	36.62		
	PLACE1005026	0.79	4.18	2.29	2.44	2.49	2.16		
	PLACE1005027	2.46	6.72	6.69	5.36	6.92	5.68		
10	PLACE1005031	1.21	1.47	3.69	5.72	7.1	4.6	*	+
	PLACE1005036	2.27	3.6	6.83	9.08	8.91	7.53	*	+
	PLACE1005041	2.5	2.84	2.84	5.05	4.41	3.25	*	+
	PLACE1005046	2.23	4.11	4.56	6.04	3.92	4.33		
15	PLACE1005047	0.23	3.19	2.6	1.43	1.23	2.1		
	PLACE1005052	4.24	8.53	6.36	8.08	8.15	8.1		
	PLACE1005055	2.54	7.45	4.66	7.2	6.45	5.62		
20	PLACE1005066	4.33	8.26	7.58	12.9	14.14	16.49	**	+
	PLACE1005077	1.17	0.68	1.2	2.1	2.43	1.54	*	+
	PLACE1005085	1.41	1.97	3.06	3.34	4.14	3.45		
	PLACE1005086	1.93	3.77	5.17	5.62	7.78	4.79		
25	PLACE1005088	24.66	32.47	46.03	43.45	31.47	27.46		
	PLACE1005089	1.57	4.78	3.15	2.52	3.67	3.14		
	PLACE1005101	3.37	8.11	5.46	6.11	8.96	6.39		
	PLACE1005102	2.56	7.14	5.01	4.11	5.51	3.8		
30	PLACE1005108	2	6.08	5.87	6.58	6.19	5.09		
	PLACE1005110	1.34	1.89	3.08	1.75	2.75	2.15		
	PLACE1005111	1.31	1.34	1.23	1.45	2.17	1.54		
35	PLACE1005123	26.23	26.21	47.58	34.26	49.34	34.98		
	PLACE1005124	3.2	4.66	4.18	4.2	6.91	5.44		
	PLACE1005128	9.54	8.89	18.22	16.37	16.36	16.13		
	PLACE1005130	2.65	6.57	5.54	2.84	4.98	3.58		
40	PLACE1005141	6.3	9.92	11.25	16.15	20.75	18.95	**	+
	PLACE1005146	1.3	2.71	3.03	2.17	2.53	2.29		
	PLACE1005152	1.85	3.9	4.56	4.13	4.23	4.79		
45	PLACE1005157	2.66	5.19	5.3	4.38	4.09	7.01		
	PLACE1005162	2.79	3.72	9.31	6.57	7.45	7.1		
	PLACE1005170	17.34	18.92	29.76	21.38	18.18	23.73		
	PLACE1005176	0.57	5.6	1.7	2.33	2.47	1.94		
50	PLACE1005181	0.53	5.14	0.96	0.89	1.36	0.37		
	PLACE1005184	4.06	9.09	10.4	8.97	12.82	11.26		
	PLACE1005186	3.5	3.41	8.56	8.05	5.79	5.73		
	PLACE1005187	2.85	4	4.13	6.1	4.99	4.25		
55	PLACE1005189	6.12	7.71	5.34	10.84	10.65	12.22	**	+

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	PLACE1004777	2.1	7.57	2.97	3.68	4.25	4.4		
	PLACE1004793	0.83	4.58	1.37	1.9	2.06	1.25		
5	PLACE1004796	6.65	8.7	13.08	7.79	8.57	7.88		
	PLACE1004804	0.99	4.46	3.25	2.44	2.38	3.22		
	PLACE1004813	4.55	7.11	9.84	6.45	5.19	5.45		
10	PLACE1004814	7.16	11.76	17.62	15.83	11.39	10.1		
	PLACE1004815	0.7	2.81	2.43	3.12	2.61	3.44		
	PLACE1004816	1.16	2.63	2.04	2.36	2.26	1.84		
	PLACE1004824	3.25	7.37	5.27	8.1	9.13	8.85		
15	PLACE1004827	1.4	10.89	3.17	2.57	3.05	1.3		
	PLACE1004836	1.72	12.95	4.26	6.25	7.99	4.49		
	PLACE1004838	1.35	8.81	2.2	2.49	2.34	1.68		
	PLACE1004840	1.59	2.06	2.21	1.85	2.08	1.37		
20	PLACE1004842	0.86	1.98	1.89	1.98	1.78	2.33		
	PLACE1004850	0.81	2.35	1.63	1.83	2.76	2.36		
	PLACE1004868	0.81	2.97	2.04	1.62	2.23	2		
25	PLACE1004885	1.5	7.09	3.51	3.4	6.03	4.77		
	PLACE1004886	1.87	8.53	2.76	3.33	5.12	4.93		
	PLACE1004887	18.14	34.01	58.51	36.9	38.66	30.33		
	PLACE1004896	8.39	14.15	15.4	9.39	11.14	14.03		
30	PLACE1004900	1.75	2.7	6.69	5.66	5.74	4.44		
	PLACE1004902	5.42	6.25	9.27	6.39	4.2	3.6		
	PLACE1004904	1.7	4.66	2.52	6.49	4.78	3.03		
35	PLACE1004911	0.69	2.5	1.12	5.95	4.82	6.09	**	+
	PLACE1004913	3.63	5.72	7.38	4.49	5.45	4.79		
	PLACE1004918	1.3	6.69	2.05	2.39	3.24	2.64		
	PLACE1004930	2.74	8.84	5.93	10.63	16.57	14.71	*	+
40	PLACE1004934	1.14	4.3	3.34	2.97	2.03	2.5		
	PLACE1004937	2.1	4.03	4.91	2.74	3.59	2.36		
	PLACE1004949	4.32	4.98	7.67	8.53	8.48	6.04		
	PLACE1004969	0.74	1.74	1.99	1.39	2.34	1.4		
45	PLACE1004970	0.45	2.18	1.2	1.43	1.31	1.01		
	PLACE1004972	1.63	6.56	2.69	5.66	3.17	4.76		
	PLACE1004974	1.27	5.21	4.2	5.6	5.7	5.62		
50	PLACE1004975	0.59	2.84	1.11	1.94	1.98	1.24		
	PLACE1004979	1.58	4.06	4.26	4.91	5.69	4.75		
	PLACE1004982	5.66	6.45	9.74	10.03	10.67	5.65		
	PLACE1004985	1.4	1.47	1.46	2.5	2.17	1.2		
55	PLACE1005003	2.85	4.22	6	6.05	6.37	5.78		

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	PLACE1004547	5	7.61	7.82	8.66	11.2	10.28		
	PLACE1004548	1.69	6.73	4.43	6.93	8.48	6.5		
5	PLACE1004550	2.27	6.24	6.67	5.92	6.78	6.15		
	PLACE1004551	0.8	2.16	1.62	2.14	2.21	1.95		
	PLACE1004559	2.9	2.89	5.11	4.45	6.75	4.82		
10	PLACE1004562	8.67	11.27	16.07	13.01	14.38	13.34		
	PLACE1004564	1.84	5.19	4.36	4.64	5.98	4.49		
	PLACE1004604	1.69	4.21	9.88	2.49	4.34	1.97		
	PLACE1004611	2.73	5.87	4.89	3.86	4.17	3.99		
15	PLACE1004629	9.42	15.75	19.92	23.93	30.49	29.01	*	+
	PLACE1004630	16.66	20.82	35.1	16.76	23.04	19.17		
	PLACE1004637	5.03	8.82	10.34	6.61	9.17	8.29		
20	PLACE1004645	36.5	39.28	92.04	85.16	87.94	74.59		
	PLACE1004646	1.07	2.91	2.87	3.68	2.19	2.28		
	PLACE1004648	0.8	3.42	2.52	2.53	3.15	1.22		
	PLACE1004655	45.95	58.09	130.94	112.14	126.25	99.01		
25	PLACE1004658	2.4	7.34	6.31	6.64	8.37	6		
	PLACE1004664	1.26	4.83	1.3	3.02	2.65	1.89		
	PLACE1004672	2.32	6.79	8.02	6.3	7.51	7.14		
	PLACE1004674	9.4	11.97	14.7	8.3	9.75	15.25		
30	PLACE1004681	1.97	3.84	5.62	5.05	5.69	4.39		
	PLACE1004686	2.74	4.33	7.46	9.22	10.77	8.59	*	+
	PLACE1004690	10.64	13.43	19.62	21.75	19.21	31.72		
35	PLACE1004691	1.14	6.71	3.71	2.92	4.13	2.75		
	PLACE1004693	1.34	7.54	4.89	3.91	4.59	5.97		
	PLACE1004701	13.01	18.45	24.24	25.21	24.46	25.1		
	PLACE1004705	1.29	3.33	2.27	1.8	1.96	1.47		
40	PLACE1004708	37.69	46.37	80.19	41.34	39.66	50.98		
	PLACE1004716	6.37	8.81	11.08	4.22	12.55	14.26		
	PLACE1004722	1.31	3.05	2.6	2.26	3.28	2.51		
45	PLACE1004736	5.25	7.71	7.6	9.16	8.89	11.63		
	PLACE1004737	5.42	12.71	16.14	8.15	11.23	13.78		
	PLACE1004740	4.88	9.06	8.22	7.37	7.93	8.2		
	PLACE1004743	1.31	4.04	3.1	1.97	4	3.55		
50	PLACE1004751	0.98	2.89	2.88	2.75	3.74	3.06		
	PLACE1004757	3.45	4.34	10.53	8.4	9.6	7.22		
	PLACE1004761	6.41	7.32	12.59	9.99	10.44	9.72		
55	PLACE1004773	1.05	2.34	1.7	1.94	2.31	2.72		
	PLACE1004775	0.35	3.26	1.37	1.29	2.14	1.07		

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	PLACE1004336	2.87	9.6	5.36	6.88	9.21	6.71		
	PLACE1004346	0.47	7.22	2.58	1.87	2.54	1.69		
5	PLACE1004358	1.3	2.41	2.41	2.46	2.09	2.03		
	PLACE1004376	11.07	10.15	23.35	19.16	17.04	16.46		
	PLACE1004384	0.65	3.46	1.46	1.92	2.48	2.1		
	PLACE1004385	1.4	2.89	1.69	3.52	1.88	2.67		
10	PLACE1004388	1.79	5.73	4.27	3.44	4.87	3.04		
	PLACE1004405	2.16	8.39	4.42	13.36	14.48	16.74	**	+
	PLACE1004407	5.05	13.12	13.37	11.2	16.24	11.85		
15	PLACE1004424	0.37	5.78	0.85	1.7	1.81	1.58		
	PLACE1004425	1.14	1.94	3.57	3.28	3.27	3.44		
	PLACE1004427	1.96	3.31	4.56	4.67	4.22	3.24		
	PLACE1004428	0.88	2.05	2.17	2.66	2.08	2.62		
20	PLACE1004433	5.7	8.3	10.82	12.94	15.67	12.05	*	+
	PLACE1004435	0.72	4.17	1.43	1.95	1.9	2.15		
	PLACE1004437	4.05	7.68	14.2	11.07	13.01	12.37		
25	PLACE1004441	7.82	11.68	34.06	30.75	43.19	26.41		
	PLACE1004446	1.5	4.36	0.9	1.03	1.35	1.39		
	PLACE1004450	0.33	1.46	1.34	2.57	1.71	0.7		
	PLACE1004451	0.51	1.45	2.14	1.89	2.69	0.88		
30	PLACE1004456	8.22	9.7	10.97	16.68	10.4	4.18		
	PLACE1004458	3.39	4.81	3.66	7.77	7.05	8.24	**	+
	PLACE1004460	0.84	4.58	2.1	2.91	2.69	1.75		
35	PLACE1004467	5.31	6.81	10.65	7.67	10.14	10.48		
	PLACE1004471	2.65	5.93	6.64	6.79	7.34	6.14		
	PLACE1004473	1.16	4.66	3.5	3.18	3.23	3.21		
	PLACE1004475	14.03	16.41	32.49	31.09	32.51	18.17		
40	PLACE1004482	8.37	6.7	10.79	10.04	9.76	9.15		
	PLACE1004491	0.39	2.51	1.49	1.19	2.68	1.3		
	PLACE1004492	61.52	74.8	127.94	129.92	127.64	123.82		
45	PLACE1004506	10.71	14.35	14.4	8.45	11.13	10.03		
	PLACE1004507	2.9	7.37	5.09	7.15	6.87	6.18		
	PLACE1004510	2.51	6.23	6.33	6.59	7.8	8.16		
	PLACE1004516	0.98	7.36	2.12	2.79	3.78	2.22		
50	PLACE1004518	1.64	1.78	3.03	2.41	3.88	2.83		
	PLACE1004519	0.17	0.82	0.62	1.43	2.79	1.51	*	+
	PLACE1004520	6.08	8.09	10.06	7.44	9.11	2.52		
	PLACE1004530	33.19	43.86	68.13	41.86	27.72	38.09		
55	PLACE1004545	1.13	3.83	2.12	3.03	3.31	3.65		

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	PLACE1004113	1.37	3.7	3.28	2.75	3.27	1.29		
	PLACE1004114	1.12	2.55	2.23	2.84	2.49	2.93		
5	PLACE1004118	1.58	3.52	-2.09	2.74	3.64	2.5		
	PLACE1004128	4.4	6.84	5.66	6.61	8.13	9.01		
	PLACE1004130	2.25	4.83	8.35	6.86	9.26	6.73		
	PLACE1004149	3.59	7.3	9.86	10.23	13.29	10.12		
10	PLACE1004156	3.61	7.91	10.12	11.66	17.62	13.31		
	PLACE1004160	5.45	9.54	14.36	13.1	19.65	14.53		
	PLACE1004161	2.2	4.86	3.54	4.85	5.73	7.48		
15	PLACE1004166	5.61	5.81	9.91	8.3	8.66	11.2		
	PLACE1004168	3.35	4.97	3.73	4.73	6.65	6.79		
	PLACE1004170	0.78	3.28	1.93	2.98	3.42	2.76		
	PLACE1004178	0.83	5.23	2.37	2.4	2.59	2.36		
20	PLACE1004183	0.89	7.99	4.41	3.53	4.32	4.84		
	PLACE1004197	0.64	5.14	1.55	1.73	3.54	1.65		
	PLACE1004199	1.66	4.52	4.09	3.78	5.88	4.35		
25	PLACE1004203	1.8	3.57	4.17	2.43	3.62	2.83		
	PLACE1004242	3.8	5.64	11.04	8.55	8.14	8.64		
	PLACE1004249	31.4	56.31	117.88	127.93	152.54	151.22	*	+
	PLACE1004255	0.79	2.65	1.26	2.59	2.15	1.93		
30	PLACE1004256	9.06	11.68	13.63	14.66	14.18	23.37		
	PLACE1004257	2.63	7.95	6.48	7.89	8.8	8.64		
	PLACE1004258	1.87	5.21	3.13	4.59	3.15	3.11		
35	PLACE1004270	0.72	3.8	2.5	2.7	4.01	1.65		
	PLACE1004272	1.34	3.68	3.73	3.86	5.38	6.15		
	PLACE1004273	92.91	89.59	212.62	212.05	129.56	99.82		
	PLACE1004274	2.09	3.61	6.51	6.42	7.14	6.74		
40	PLACE1004277	2.3	4.4	5.76	6.45	7.7	6.04		
	PLACE1004279	0.54	3.39	2.23	3.16	2.64	2.02		
	PLACE1004282	2.43	8.25	6.62	4.22	5.56	4.49		
45	PLACE1004284	4.59	11.31	7.84	7.38	8.16	7.15		
	PLACE1004289	1.28	7.85	2.46	3.06	3.63	2.1		
	PLACE1004299	0.33	6.41	1.38	1.54	2.67	1.83		
	PLACE1004302	1.01	2.98	3.27	2.41	2.45	1.12		
50	PLACE1004305	1.11	2.09	1.9	1.9	1.82	1.78		
	PLACE1004316	2.3	4.48	5.4	6.06	3.85	4.52		
	PLACE1004322	2.49	3.41	5.25	6.35	7.14	5.75	*	+
	PLACE1004325	2.43	6.38	3.84	3.85	3.66	4		
55	PLACE1004332	1.21	7.18	2.8	3.72	3.46	3.01		

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	PLACE1003768	0.36	1.14	1.36	1.45	2.75	1.07		
	PLACE1003771	1.28	1.94	2.07	1.84	3.13	1.43		
5	PLACE1003772	34.15	38.19	-97.86	62.42	64.06	52.43		
	PLACE1003783	1.48	3.02	2.22	18.65	19.53	16.61	**	+
	PLACE1003784	0.69	3.92	0.87	2.09	2.19	2.68		
10	PLACE1003788	0.4	4.92	1.06	1.85	1.71	0.32		
	PLACE1003795	1.01	4.1	3.57	4.73	4.38	3.54		
	PLACE1003827	13.83	20.46	20.72	22.48	30.84	25.92		
	PLACE1003833	0.98	1.49	3.9	3.65	4.33	3.31		
15	PLACE1003839	22.55	19.18	52.95	50.39	56.11	43.86		
	PLACE1003845	6.09	6.88	11.72	24.98	19.99	10.6		
	PLACE1003850	3.16	4.84	7.19	5.45	5.95	6.39		
20	PLACE1003852	0.25	3.36	1.09	0.99	1.58	1.05		
	PLACE1003858	1.34	4.42	1.99	2.04	1.85	2.92		
	PLACE1003861	0.95	4.51	1.63	2.98	2.78	1.73		
	PLACE1003864	0.94	5.5	2.74	2.88	3.86	2.69		
25	PLACE1003870	3.84	3.4	13.2	10.71	16.12	12.03		
	PLACE1003885	1.33	1.42	1.59	3.07	3.76	2.38	*	+
	PLACE1003886	4.56	6.01	5.75	9.27	6.87	4.3		
	PLACE1003888	0.75	3.79	1.96	2.87	3.42	2.68		
30	PLACE1003892	4.93	6.91	20.79	17.21	22.33	14.87		
	PLACE1003900	2.27	5.92	7.17	6.34	9.75	6.6		
	PLACE1003902	1.91	7.39	5.25	5.43	8.07	6.29		
35	PLACE1003903	0.42	5.07	2.94	2.55	3.5	2.59		
	PLACE1003915	8.15	7.04	10.78	8.31	9.79	9.84		
	PLACE1003918	1.88	2.45	4.75	3.47	6.26	3.75		
	PLACE1003923	2.06	3.73	5.63	2.7	5.54	3.32		
40	PLACE1003932	3.99	5.16	5.47	4.06	7.58	5.09		
	PLACE1003936	1.02	3.82	2.81	3.63	2.42	2.78		
	PLACE1003966	3.11	7.43	7.76	4.89	7.32	4.21		
	PLACE1003968	1.68	5.68	5.94	3.33	4.26	4.57		
45	PLACE1004018	25.49	33.73	48.16	32.56	40.53	28.62		
	PLACE1004020	8.91	10.18	13.26	11.42	18.03	24.15		
	PLACE1004028	0.41	2.55	1.3	1.38	1.23	1.91		
50	PLACE1004034	3.56	4.53	5.22	8.42	9.03	13.04	*	+
	PLACE1004042	17.25	20.19	68.35	79.51	93.32	79.44	*	+
	PLACE1004078	1.14	4.1	3.3	4.95	6.51	4.45		
	PLACE1004103	5.54	10.93	13.98	13.77	15.09	14.14		
55	PLACE1004104	5.94	12.29	27.78	24.18	34.98	29.71		

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	PLACE1003573	0.98	2.43	1.19	2.16	1.87	1.48		
	PLACE1003575	2.16	3.09	3.44	3.34	3.5	4.15		
5	PLACE1003583	0.97	3.45	-3.34	2.29	4.16	2.23		
	PLACE1003584	1.23	4.46	4.01	3.56	3.8	2.65		
	PLACE1003592	4.4	8.48	9.48	7.11	10.48	9.53		
10	PLACE1003593	0.84	5.55	2.4	2.23	3.02	1.55		
	PLACE1003594	4.24	6.76	6.01	5.78	6.68	7.36		
	PLACE1003596	13.77	11.31	22.53	15.68	17.69	8.25		
	PLACE1003598	2.83	3.63	5.02	4.89	3.7	2.95		
15	PLACE1003602	1.8	4.24	6.36	4.37	3.54	2.68		
	PLACE1003605	17.43	21.72	45.86	94.65	95.6	91.55	**	+
	PLACE1003611	2.34	5.18	6.07	5.65	7.14	6		
	PLACE1003618	0.67	7.39	2.09	1.58	2.4	1.32		
20	PLACE1003625	1.78	10.41	2.75	3.33	5.48	2.8		
	PLACE1003626	8.77	15.99	17.14	10.87	14.46	13.19		
	PLACE1003630	1.8	2.57	5.8	7.05	4.67	5.86		
25	PLACE1003635	2.15	1.83	3.19	2.96	2.82	2.44		
	PLACE1003638	1.3	2.58	3	5.21	4.19	3.13		
	PLACE1003644	4.01	5.7	7.25	7.81	9.13	8.76	*	+
	PLACE1003654	2.56	6.14	4.04	3.54	6.96	6.4		
30	PLACE1003656	2.69	7.79	6.12	6.54	5.63	4.77		
	PLACE1003660	0.26	9.54	3.5	3.08	4.92	4.11		
	PLACE1003669	2.43	9.05	3.67	2.59	4.26	3.08		
35	PLACE1003670	5.37	5.7	9.44	11.01	8.26	8.76		
	PLACE1003671	1.66	1.22	3.57	3.11	2.57	1.57		
	PLACE1003697	7.27	7.99	9.8	8.23	6.06	6.42		
	PLACE1003704	3.12	3.97	5.17	5.96	7.25	5.97	*	+
40	PLACE1003709	0.89	2.63	0.8	1.19	1.24	2.44		
	PLACE1003711	0.74	5.48	1.35	1.87	1.8	1.39		
	PLACE1003723	1.07	6.99	4.7	4.2	5.31	4.16		
45	PLACE1003724	3.31	10.74	9.1	9.11	11.79	10.49		
	PLACE1003737	2.14	2.21	4.72	3.35	3.1	3.29		
	PLACE1003738	1.06	1.94	3.13	3.96	3.92	3.41		
	PLACE1003742	2.25	3.58	5.71	6.81	6.18	2.85		
50	PLACE1003744	6.13	8.86	14.6	16.21	17.96	19.13	*	+
	PLACE1003758	0.85	4.55	0.96	2	1.46	1.16		
	PLACE1003760	13.44	18.68	27.23	31.82	20.52	22.79		
	PLACE1003762	1.45	4.97	3.7	3.77	3.78	3.49		
55	PLACE1003765	1.18	5.23	3.45	2.01	3.1	2.11		

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	PLACE1003353	19.67	16.04	40.09	37.78	42.55	40.26		
	PLACE1003361	1.82	3.64	3.72	5.85	5.31	4.6	*	+
5	PLACE1003366	1.45	4.35	3.63	3.22	3.33	2.97		
	PLACE1003369	2.75	4.51	3.49	3.29	3.25	4.62		
	PLACE1003372	2.08	5.73	2.68	5.45	4.72	3.39		
10	PLACE1003373	2.85	7.37	6.62	8.8	11.81	9.89		
	PLACE1003375	1.42	4.91	1.92	2.59	2.95	3.21		
	PLACE1003378	0.94	0.94	0.98	0.7	1.66	1.08		
	PLACE1003383	0.87	1.55	2.33	1.59	3.15	1.57		
15	PLACE1003394	10.55	12.49	24.08	11.75	22.99	17.27		
	PLACE1003401	0.79	3.91	1.34	1.03	2.13	1.04		
	PLACE1003405	1.5	3.97	2.22	2.54	2.46	2.04		
20	PLACE1003407	2.39	6.06	5.16	3.96	6.3	4.02		
	PLACE1003420	3.26	7.69	6.19	6.8	10.92	8.7		
	PLACE1003428	0.63	3.3	2.62	2.07	2.94	1.96		
	PLACE1003432	6.14	5.81	8.2	6.64	7.05	5.42		
25	PLACE1003438	0.45	2.66	0.93	2.41	2.34	1.99		
	PLACE1003452	1.87	5.02	5.08	4.53	3.43	3.84		
	PLACE1003454	2.49	5.59	7.34	7.31	6.95	5.61		
	PLACE1003455	2.58	4.26	2.35	2.97	3.01	3.17		
30	PLACE1003456	3.22	7.74	8.62	6.9	7.2	7.79		
	PLACE1003460	6.39	13.35	14.87	13.02	16.76	12.86		
	PLACE1003478	1.15	1.71	0.86	2.33	2.07	1.24		
35	PLACE1003484	12.06	12.21	45.33	28.12	31.5	34.2		
	PLACE1003493	1.61	4.72	4.9	3.84	5.96	5.08		
	PLACE1003503	85.45	87.35	107.79	115.17	111.85	172.81		
	PLACE1003505	1.99	6.77	4.78	7.44	6.63	8.87		
40	PLACE1003516	0.86	6.78	2.7	2.8	3.95	2.39		
	PLACE1003519	17.58	26.29	50.41	45.77	36.97	58.75		
	PLACE1003520	14.18	25.48	35.96	25.73	31.4	32.19		
45	PLACE1003521	2.71	3.64	4.93	5.97	4.71	7.4		
	PLACE1003525	8.45	11.81	45.05	33.94	43.71	36.88		
	PLACE1003528	39.18	44.68	136.4	106.04	122.76	127.22		
	PLACE1003529	1.46	4.26	3.29	2.4	3.94	3.83		
50	PLACE1003537	4.41	9.05	11.05	11.36	13.77	13.3		
	PLACE1003549	1.1	5.02	4.59	5.61	6.01	5.93		
	PLACE1003553	1.6	5.89	3.88	4.02	4.17	4.23		
	PLACE1003566	5.93	9.8	17.51	13.03	19.09	14.58		
55	PLACE1003568	3.01	2.71	5.76	5.69	4.43	4.04		

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	PLACE1003085	3.91	6.19	5.6	9.46	5.89	3.33		
	PLACE1003092	3.94	4.87	6.25	7	5.6	6.17		
5	PLACE1003097	0.37	3.06	-1.44	2.12	1.88	1.63		
	PLACE1003100	1.65	7.1	4.2	3.88	4.74	4.29		
	PLACE1003108	1.26	10.37	2.91	3.32	4.44	2.39		
10	PLACE1003115	11.39	18.3	58.59	73.64	99.24	69.1	*	+
	PLACE1003120	3.1	3.08	9.71	11.34	8.32	10.19		
	PLACE1003135	0.72	2.04	1.09	1.56	2.89	1.08		
	PLACE1003136	3.95	5.82	6.05	9.03	6.55	7.34		
15	PLACE1003141	2.04	2.97	2.1	1.97	2.49	1.8		
	PLACE1003145	1.21	4.17	2.52	6.24	6.88	7.67	*	+
	PLACE1003147	2.87	7.85	5.71	5.02	5.25	6.28		
20	PLACE1003153	0.54	7.63	2.14	1.66	3.2	1.82		
	PLACE1003163	6.09	13.55	8.19	8.39	14.09	12.26		
	PLACE1003172	23.21	21.74	44.19	47.78	43.17	39.52		
	PLACE1003174	2.31	2.49	3.75	4.3	3.55	1.68		
25	PLACE1003176	0.47	2	1.89	2.88	1.27	1.46		
	PLACE1003181	1.72	4.19	2.72	2.5	2.76	2.36		
	PLACE1003184	0.76	3.92	1.53	1.91	1.49	2.06		
30	PLACE1003190	2.39	9.81	8.67	10.73	7.98	10.34		
	PLACE1003200	0.29	4.48	1.84	0.72	1.92	1.16		
	PLACE1003205	3.94	7.07	9.68	6.82	10.38	7.2		
	PLACE1003209	1.43	2.18	2.62	2.28	1.82	2.89		
35	PLACE1003214	0.83	1.3	2	2.15	2.44	1.81		
	PLACE1003229	2.08	2.78	2.9	2.78	3.35	3.48		
	PLACE1003238	0.46	2.34	1.24	1.35	2	0.75		
40	PLACE1003249	1.87	5.04	4.7	7.33	7.56	5.89		
	PLACE1003256	3.47	7.69	7.94	8.82	7.68	6.08		
	PLACE1003258	1.03	3.81	3.48	2.42	2.19	0.87		
	PLACE1003279	3.09	7.19	9.02	11.15	13.56	11.58	*	+
45	PLACE1003294	0.95	1.54	1.59	1.57	1.25	2.64		
	PLACE1003296	1.49	2.6	2.45	2.59	3.11	2.4		
	PLACE1003297	7.52	10.15	31.88	23.01	23.49	19.3		
	PLACE1003302	3.92	5.16	6.99	5.8	4.72	5.47		
50	PLACE1003334	1.51	4.41	1.91	2.4	3.59	3.09		
	PLACE1003337	13.69	16.3	29.83	28.53	34.27	25.76		
	PLACE1003342	1.05	4.07	1.89	2.15	2.35	1.97		
	PLACE1003343	1.07	4.98	1.61	2.02	2.75	2.12		
55	PLACE1003344	6.25	5.33	12.83	11.18	11.35	11.98		

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	PLACE1002775	4.33	4.71	9.15	7.05	7.08	8.67		
	PLACE1002780	185.63	218.72	325.36	272.21	244.38	305.38		
5	PLACE1002782	0.4	3.76	1.1	1.62	1.69	1.14		
	PLACE1002794	1.5	6.71	3.27	2.26	4.59	4.36		
	PLACE1002795	1.92	6.45	0.81	2.37	3.63	2.77		
	PLACE1002811	0.6	1.57	1.34	1.9	1.46	1.16		
10	PLACE1002815	6.39	7	10.49	7.24	3.16	9.21		
	PLACE1002816	8.5	9.72	9.05	7.22	8.2	7.97	*	-
	PLACE1002822	0.58	2.51	2.06	2.2	2.87	1.94		
15	PLACE1002833	9.98	15.11	16.78	15.42	15.99	16.56		
	PLACE1002834	3.2	8.08	6.57	6.23	6.79	8.41		
	PLACE1002835	0.62	4.14	1.72	0.85	1.83	1.79		
	PLACE1002839	1.13	5.75	2.72	2.89	4.72	2.81		
20	PLACE1002851	1.52	1.87	1.41	1.98	2.15	2.7		
	PLACE1002853	4.18	6.23	9.15	6.26	5.6	7		
	PLACE1002881	3.42	5.2	11.04	8.35	11.57	10.26		
25	PLACE1002901	9.66	12.66	20.09	24.14	25.51	30.67	*	+
	PLACE1002904	0.89	7.35	1.41	1.98	1.95	3.09		
	PLACE1002905	1.36	5.46	3.26	4.04	4.46	3.4		
	PLACE1002908	1.6	5.19	3.18	3.84	5.27	3.81		
30	PLACE1002911	3.91	6.96	6.9	4.66	7.89	6.75		
	PLACE1002941	1.57	2.2	2.48	3.94	2.02	2.31		
	PLACE1002950	9.59	9.15	14.74	5.31	8.02	14.51		
35	PLACE1002955	47.83	40.69	72.7	82.17	62.5	84.64		
	PLACE1002958	19.36	26.92	35.27	35.6	35.35	59.02		
	PLACE1002962	1.03	4.03	2.2	1.41	2.63	1.67		
	PLACE1002967	1.34	4.83	3.19	4.37	3.52	2.81		
40	PLACE1002968	1.2	5.14	2.7	2.55	3.05	1.81		
	PLACE1002976	8.94	12.08	24.23	24.5	36.89	30.05		
	PLACE1002991	2.68	3.05	6.66	3.49	4.56	3.6		
	PLACE1002993	2.72	3.86	5.52	8.21	6.92	5.56		
45	PLACE1002996	2.02	3.03	3.43	5.54	3.52	3.01		
	PLACE1003010	1.91	3.69	4.27	4.31	3.86	3.32		
	PLACE1003025	2.85	7.01	6.1	8.57	11.37	10.11	*	+
50	PLACE1003027	5.02	13.08	9.31	8.55	12.45	12.76		
	PLACE1003044	1.95	8.24	2.61	3.64	4.16	2.74		
	PLACE1003045	1.41	7.75	1.77	1.88	2.64	1.01		
	PLACE1003052	2.19	3.16	5.74	4.44	3.6	1.99		
55	PLACE1003083	1.59	3.04	3.23	3.06	1.61	2.25		

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	PLACE1002446	4.64	8.42	5.95	10.27	8.88	11.99	*	+
	PLACE1002447	1.26	6.06	2.05	3.92	3.14	4.32		
5	PLACE1002450	1.19	5.92	-3.24	4.32	4.21	5.05		
	PLACE1002462	0.81	4.02	2.94	3.51	2.99	3.13		
	PLACE1002465	0.96	4.69	2.2	2.69	4.31	2.24		
	PLACE1002474	1.61	2.26	3.23	3.85	4.41	3.6	*	+
10	PLACE1002477	11.11	14.51	32.39	44.06	41.42	26.68		
	PLACE1002493	3.39	4.1	10.1	14.39	16.66	9.49		
	PLACE1002497	0.68	2.81	0.67	1.45	0.93	0.99		
15	PLACE1002499	2.12	4.73	3	5.98	6.44	5.28	*	+
	PLACE1002500	2.61	6.52	7.36	7.58	10.45	7.25		
	PLACE1002514	0.3	4.49	1.84	1.74	2.47	1.75		
	PLACE1002518	2.86	7.65	6.9	5.62	7.55	4.67		
20	PLACE1002529	1.14	1.56	2.21	3.19	3.4	1.44		
	PLACE1002532	1.31	1.82	3.18	5.75	4.94	5.59	**	+
	PLACE1002536	3.59	3.75	3.44	5.84	6.07	3.85		
25	PLACE1002537	1.63	4.06	2.7	2.69	4.07	3.08		
	PLACE1002539	1.86	5.68	2.75	4.53	5.29	4.78		
	PLACE1002547	6.09	8.06	7.3	12.32	11.02	11.26	**	+
	PLACE1002571	2.84	6.85	5.19	6.84	8.65	6.23		
30	PLACE1002578	3.57	8.34	8.35	11.11	12.19	8.11		
	PLACE1002583	1.33	1.61	2.32	3.18	4.02	2.46		
	PLACE1002591	0.82	1.62	2.34	3.25	4.43	1.92		
35	PLACE1002598	6.56	10.95	12.39	11.93	9.04	7.74		
	PLACE1002604	1.73	3.57	2.69	3.75	5.38	3.51		
	PLACE1002612	2.89	8.47	5.95	11.25	10.88	8.06		
	PLACE1002625	1.25	4.79	3.18	2.7	3.25	1.82		
40	PLACE1002638	2.94	8.01	6.66	7.78	6.81	7.29		
	PLACE1002655	1.39	6.51	5.57	7.19	7.62	6.46		
	PLACE1002665	4.57	3.88	5.4	7.47	12.16	10.75	*	+
	PLACE1002685	0.58	1.12	1.3	0.67	2.43	0.98		
45	PLACE1002692	7.42	8.56	16.7	19.27	22.67	16.29		
	PLACE1002714	1.8	3	2.11	2.43	3.14	2.24		
	PLACE1002721	2.94	4.37	3.88	5.88	7.1	4.28		
50	PLACE1002722	0.92	5.42	1.97	1.37	3.28	1.85		
	PLACE1002726	1.6	6.24	3.66	4.6	5.7	5.26		
	PLACE1002756	1.57	4.5	7.04	5.92	9.63	7.78		
	PLACE1002768	1.05	3.72	2.16	2.1	2.34	1.71		
55	PLACE1002772	0.54	2.15	1.32	2.49	2.86	2.3		

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	PLACE1002150	1.54	8.26	4.25	3.23	5.36	4.12		
	PLACE1002153	1.6	5.75	2.58	3.47	5.76	4.48		
5	PLACE1002157	0.87	2.96	1.72	1.76	3.28	3.73		
	PLACE1002163	2.13	4.67	4.55	6.21	8.81	6.03		
	PLACE1002168	2.39	4.04	4.42	4.68	6.28	3.22		
	PLACE1002170	2.73	3.53	7.35	4.89	5.33	3.74		
10	PLACE1002171	3.09	6.11	10.02	18.44	14.63	13.93	*	+
	PLACE1002180	3.16	6.23	6.77	4.63	8.54	8.39		
	PLACE1002184	9.2	15.58	18.42	124.63	250.27	333.14	*	+
15	PLACE1002200	1.35	5.67	2.38	2.97	2.89	2.26		
	PLACE1002205	3.3	6.47	18.07	17.08	18.38	14.61		
	PLACE1002213	2.2	4.16	4.93	5.52	8.03	6.03		
	PLACE1002219	1.05	1.91	2.23	3.33	3.53	1.96		
20	PLACE1002227	0.68	2.65	1.7	1.5	3.03	1.67		
	PLACE1002253	0.32	2.67	1.28	1.47	0.76	0.43		
	PLACE1002256	1.16	4.78	3.31	3.54	3.01	4.79		
25	PLACE1002259	1.46	5.69	4.48	3.22	2.98	2.31		
	PLACE1002285	1.16	10.74	2.29	1.55	2.38	1.24		
	PLACE1002301	9.42	17.5	14.68	12.7	10.48	11.7		
	PLACE1002310	4.28	10.16	9.86	8.82	7.87	9.94		
30	PLACE1002311	1.84	2.94	3.87	2.96	2.87	2.03		
	PLACE1002319	2.31	2.64	2.94	3.21	3.23	3.92	*	+
	PLACE1002329	0.56	2.54	2.5	4.07	3.58	3.07		
35	PLACE1002333	1.34	3.1	1.96	1.22	2.44	2		
	PLACE1002342	4.19	9.04	9.44	5.06	8.52	8.17		
	PLACE1002343	0.49	6.98	2.94	2.08	1.9	2.52		
	PLACE1002355	1.31	9.39	2.36	3.33	4.35	2.63		
40	PLACE1002358	1.15	7.94	3.3	2.6	2.65	2.13		
	PLACE1002359	1.91	2.17	3.47	4.7	3.91	3.42		
	PLACE1002374	29.69	28.18	54.19	53.9	34.73	36.14		
	PLACE1002376	3.58	5.91	7.86	6.23	6.82	6.56		
45	PLACE1002379	6.24	7.66	6.63	10.13	9.68	10.9	**	+
	PLACE1002386	0.86	5.32	1.35	1.87	2.05	1.51		
	PLACE1002395	3.69	9.97	17.13	16.43	20.62	16.16		
50	PLACE1002399	2.38	11.09	3.42	5.31	10.38	7.39		
	PLACE1002407	1.09	5.22	2.31	2.3	4.01	3.66		
	PLACE1002433	1.63	2.17	2.97	2.96	4.35	3.66		
	PLACE1002437	0.79	1.4	1.47	1.41	3.28	1.35		
55	PLACE1002438	0.74	2.38	1.96	1.8	2.43	3.38		

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	PLACE1001845	3.62	8.38	7.39	7.88	7.55	9.93		
	PLACE1001858	2.56	6.58	3.52	5.26	7.48	6.05		
5	PLACE1001869	3.13	7.15	-4.85	6.09	6.46	5.66		
	PLACE1001890	11.74	11.92	21.45	173.44	255.31	125.13	*	+
	PLACE1001897	9.19	13.85	16.44	22.22	23.13	12.95		
10	PLACE1001902	10.13	12.6	21.53	22.74	27.67	12.77		
	PLACE1001904	1.38	3.72	1.51	2.45	2.53	2.35		
	PLACE1001907	3.36	6.76	5.71	7.67	5.67	5.59		
	PLACE1001910	83.6	82.16	135.34	301.29	325.42	244.59	**	+
15	PLACE1001912	1.53	6.6	3.36	5.54	5.48	4.85		
	PLACE1001918	17.31	22.95	30.16	31.14	40.44	40.02	*	+
	PLACE1001920	2.07	3.51	5.43	11.97	13.8	11.4	**	+
	PLACE1001928	3.06	2.96	4.67	5.29	9.7	5.31		
20	PLACE1001930	1.17	3.92	2.2	2.9	4.73	3.22		
	PLACE1001949	1.16	3.67	1.78	3.84	4.24	3.18		
	PLACE1001959	1.36	4.7	3.16	2.63	3.17	2.26		
25	PLACE1001969	2.09	7.83	7.21	6.56	10.73	6.57		
	PLACE1001974	7.39	11.98	11.87	11.43	16.09	16.06		
	PLACE1001981	0.77	4.38	3.22	1.77	3.88	2.36		
	PLACE1001983	3.81	4.12	5.32	5.92	6.16	5.72	*	+
30	PLACE1001989	2.34	4.15	5.02	4.37	5.91	3.72		
	PLACE1002004	3.07	4.06	8.05	9.22	9.69	7.18		
	PLACE1002008	8.4	11.76	17	23.36	22.19	22.42	*	+
35	PLACE1002015	26.96	30.92	67.62	105.75	88.42	94.15	*	+
	PLACE1002044	3.79	8.07	5.86	4.64	6.39	6.4		
	PLACE1002046	1.78	5.68	1.9	4.3	5.79	4.97		
	PLACE1002052	1.09	4.98	2.26	1.38	2.41	2.32		
40	PLACE1002066	4.79	6.3	8.29	10.24	10.77	9.93	*	+
	PLACE1002072	2.55	3.91	4.86	6	5.48	6	*	+
	PLACE1002073	0.51	2.83	2.29	2.35	4.06	2.91		
	PLACE1002080	1.81	6.49	6.13	4.76	6.82	5.72		
45	PLACE1002081	1.66	6.13	4.06	3.74	4.86	4.3		
	PLACE1002090	7.74	16.55	13.87	12.53	14.4	19.41		
	PLACE1002095	2.97	6.22	8.45	10.01	10.18	11.32	*	+
50	PLACE1002102	4.26	8.56	8.81	9.47	9.56	10.67		
	PLACE1002109	2.57	5.08	3.81	4.66	6.17	6.32		
	PLACE1002115	1.75	3.57	2.48	2.78	4.26	2.59		
	PLACE1002119	15.65	15.3	35.78	37.28	32.59	38.23		
55	PLACE1002140	5.25	8.45	14.05	19.93	17.14	17.19	*	+

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	PLACE1001629	0.78	6.77	2.24	3.62	3.36	3.52		
	PLACE1001632	1.66	8.26	4.04	4.3	4.14	4.37		
5	PLACE1001634	7.4	9.92	39.12	23.85	32.41	18.38		
	PLACE1001637	0.84	2.16	1.25	1.41	2.4	1.1		
	PLACE1001640	1.33	3.27	4.66	2.68	4.85	4.49		
10	PLACE1001655	0.83	2.93	2.06	2.82	2.14	2.02		
	PLACE1001672	1.84	7.04	4.01	3.3	4.41	4.09		
	PLACE1001676	1.38	8.49	3.54	4.63	4.77	3.85		
	PLACE1001683	12.79	23.62	24.61	25.33	30.22	27.13		
15	PLACE1001691	3.41	12.29	6.72	9.03	8.96	9.83		
	PLACE1001692	1.47	2.96	5.25	5.87	5.6	5.13		
	PLACE1001705	3.02	3.75	9.88	10.06	9.21	8.32		
	PLACE1001716	1.68	3	2.61	2.24	3.79	3.58		
20	PLACE1001720	1.49	2.62	2.21	1.56	2.45	1.71		
	PLACE1001728	1.43	6.19	4.24	1.96	2.04	2.51		
	PLACE1001729	2.12	8.13	4.44	3.8	4.52	4.36		
25	PLACE1001739	2.61	9.55	4.04	4.95	7.24	6.16		
	PLACE1001740	0.92	5.36	2.09	1.92	2.1	1.69		
	PLACE1001745	1.15	0.98	3.22	1.87	2.48	2.31		
	PLACE1001746	1.04	2.25	2.55	4.64	3.4	2.82		
30	PLACE1001748	4.74	7.01	8.18	8.19	6.58	5.96		
	PLACE1001753	2.06	3.54	3.29	7.44	5.57	5.82	*	+
	PLACE1001756	5.6	11.31	38.07	31.78	44.99	35.99		
35	PLACE1001760	6.54	12.23	12.85	16.36	16.96	16.66	*	+
	PLACE1001767	11.26	14.98	59.72	45.37	61.46	45.39		
	PLACE1001771	1.96	6.64	4.03	4.32	5.22	4.54		
	PLACE1001775	2.23	2.81	6.72	5.1	3.11	4.79		
40	PLACE1001777	83.34	145.9	190.82	142.92	71.27	59.69		
	PLACE1001781	1.9	3.86	4.91	8.72	3.39	2.3		
	PLACE1001783	0.76	3.21	2.06	4.84	2.09	1.54		
	PLACE1001786	1.77	6.61	2.72	3.7	3.32	2.6		
45	PLACE1001788	5.16	9.07	7.14	10.52	8.74	9.32		
	PLACE1001795	1.92	4.4	4.82	5.42	4.61	5.42		
	PLACE1001799	0.69	3.62	2.11	1.86	2.83	1.97		
50	PLACE1001810	0.89	1.52	1.76	2.73	3.91	1.73		
	PLACE1001817	5.53	6.12	10.88	10.56	9.4	6.38		
	PLACE1001821	4.68	6.07	7.11	8.37	9.92	4.99		
	PLACE1001836	0.91	3.12	2.38	2.69	4.12	2.63		
55	PLACE1001844	1.55	5.1	3.48	4.42	4.36	4.09		

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	PLACE1001377	3.02	4.97	5.1	7.91	7.28	5.83	*	+
	PLACE1001383	2.31	4.13	3.53	2.62	5.5	5.72		
5	PLACE1001384	1.81	3.23	2.89	2.05	3.43	3.15		
	PLACE1001387	1.65	3.64	3.7	3.03	4.83	3		
	PLACE1001395	3.72	6.64	6.54	7.01	7.61	7.73		
	PLACE1001399	3.71	6.58	9.31	7.37	8.61	9.22		
10	PLACE1001401	0.83	5.25	2.33	1.55	1.76	0.87		
	PLACE1001407	11.65	21.8	24.47	22.63	18.09	26.24		
	PLACE1001412	1.6	4.98	4.53	4.08	4.42	3.83		
15	PLACE1001414	2.3	3.02	5.86	7.57	5.13	6.83		
	PLACE1001416	2.99	4.71	3.29	5.62	4.04	7.08		
	PLACE1001433	33.62	33.05	51.64	49.1	58.33	55.88		
	PLACE1001440	1.95	3.99	3.96	3.6	3.53	2.1		
20	PLACE1001456	1.64	5.5	4.26	4.15	4.87	4.49		
	PLACE1001464	32.76	28.05	47.41	53.22	68.42	61.32	*	+
	PLACE1001468	0.85	5.04	1.17	1.56	2.55	2.27		
25	PLACE1001484	1.31	4.85	2.96	4.25	5.8	3.04		
	PLACE1001500	0.92	2.22	2.14	2.72	3.34	3.26	*	+
	PLACE1001502	1.36	3.6	3.9	3.54	5.9	4.54		
	PLACE1001503	1.7	4.58	6.72	7.47	8.2	8.05		
30	PLACE1001505	6.34	14.13	16.16	39.97	27.14	46.65	*	+
	PLACE1001513	4.09	10.82	8.17	5.87	8.53	14.61		
	PLACE1001516	0.61	4.33	1.33	1.71	3.49	1.99		
35	PLACE1001517	5.56	8.58	14.77	14.14	14.96	14.28		
	PLACE1001523	12.83	14.09	20.42	22.79	19.74	32.9		
	PLACE1001526	5.12	4.89	8.42	9.51	9.11	6.89		
	PLACE1001534	2.12	5.12	3.58	3.62	5.55	3.99		
40	PLACE1001536	0.61	2.5	1.52	2.11	3.2	1.9		
	PLACE1001545	17.97	23.9	38.46	33.78	45.13	66.08		
	PLACE1001551	2.55	6.26	6.15	4.72	6.59	6.71		
45	PLACE1001564	1.37	4.87	2.88	4.01	3.57	3.7		
	PLACE1001570	2.62	5.95	4.18	2.19	3.82	4.32		
	PLACE1001571	2.04	4.51	6.07	5.69	6.27	5.81		
	PLACE1001595	4.73	4.64	10.04	11.6	8.27	5.28		
50	PLACE1001602	7.23	8.39	18.65	20.38	18.68	19.71		
	PLACE1001603	2.01	3.83	5.37	6.86	5.86	4.56		
	PLACE1001608	3.44	7.22	5.9	5.82	7.73	8.7		
	PLACE1001610	3.77	8.4	8.22	9.26	9.49	9.85		
55	PLACE1001611	1.94	7.34	3.65	2.28	3.85	1.88		

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	PLACE1001092	2.31	2.47	4.68	5.8	4.18	3.9		
	PLACE1001098	0.93	2.62	4.53	5.49	4.05	3.17		
5	PLACE1001100	1.31	2.58	2.48	4.27	4.17	3.37	*	+
	PLACE1001104	1.95	4.09	4.54	3.39	3.42	4.47		
	PLACE1001114	1.56	6.54	4.33	5.17	3.78	3.27		
10	PLACE1001118	2.52	5.77	6.12	6.21	6.14	5.1		
	PLACE1001123	2.86	5.3	7.53	7.08	8.51	7.63		
	PLACE1001136	1.58	4.39	5.13	5.29	5.95	5.85		
	PLACE1001144	6.27	5.67	13.43	10.34	11	10.08		
15	PLACE1001147	2.11	2.98	6.03	6.13	5.15	4.83		
	PLACE1001148	1.72	1.31	1.89	1.76	2.39	2.31		
	PLACE1001159	0.86	2.37	2.05	2.27	3.73	1.43		
20	PLACE1001168	8.87	14.52	15.09	25.46	23.18	30.79	*	+
	PLACE1001171	0.69	3.89	1.23	2.53	1.42	1.53		
	PLACE1001183	0.24	3.61	1.81	1.57	2.78	1.38		
	PLACE1001185	3.13	7.43	3.76	5	6.4	5.64		
25	PLACE1001201	1.77	2.8	3.29	6.32	6.94	6.32	**	+
	PLACE1001229	7.51	8.56	12.64	15.24	11.45	10.42		
	PLACE1001231	1.83	2.73	3.07	4.09	5.1	2.3		
	PLACE1001238	1.52	4.35	3.74	3.65	4.52	4.57		
30	PLACE1001241	1.63	5.58	2.92	5.73	8.13	7.04		
	PLACE1001242	22.28	29.54	30.28	46.43	48.89	62.65	*	+
	PLACE1001247	2.43	7.02	4.07	5.03	5.91	4.52		
35	PLACE1001250	1.01	5.36	3.61	4.68	4.39	4.81		
	PLACE1001257	2.99	3.06	7.06	7.89	9.21	7.69		
	PLACE1001272	3.19	4.27	5.68	7.13	6.43	5.14		
	PLACE1001279	0.96	3.12	2.74	3.08	3.81	3.29		
40	PLACE1001280	1.08	4.75	2.68	4.98	4.45	2.86		
	PLACE1001294	1.91	7.23	6.91	4.88	5.57	6.18		
	PLACE1001295	4.16	9.94	7.53	8.55	11.85	8.43		
45	PLACE1001300	2.46	7.9	4.31	4.65	14.73	4.95		
	PLACE1001304	3	8.27	10.47	8.57	10.81	10.64		
	PLACE1001311	3.95	3.34	5.67	6.85	9.14	7.6	*	+
	PLACE1001323	2.17	2.95	5.12	5.66	8.43	5.5		
50	PLACE1001325	0.88	1.95	3.71	2.84	3.56	3.27		
	PLACE1001340	5.18	6.99	9.8	8.69	12.02	10.48		
	PLACE1001344	1.52	3.49	1.77	2.34	2.06	1.75		
	PLACE1001351	3.23	6.39	8.39	6.4	8.62	6.1		
55	PLACE1001366	1	4.49	4.02	4.19	4.6	3.72		

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	PLACE1000838	16	15.77	23.73	13.88	15.6	15.65		
	PLACE1000841	1.22	3.78	3.31	3.97	10.65	3.77		
5	PLACE1000843	2.14	6.2	5.68	5.79	7.7	5.38		
	PLACE1000849	2.79	8.82	6.72	7.24	6.78	10.02		
	PLACE1000856	2.01	5.3	3.59	3.42	4.79	4.19		
10	PLACE1000863	5.2	7.58	9.56	8.97	12.34	11.53		
	PLACE1000876	3.65	7.6	6.02	6.7	9.95	9.06		
	PLACE1000899	1.36	2.24	3.12	4.12	5.14	4.22	*	+
	PLACE1000907	4.82	5.53	9.59	6.77	8.44	5.83		
15	PLACE1000909	1.18	3.31	2.45	3.65	3.88	3.44		
	PLACE1000912	0.42	4.55	1.77	1.76	2.72	1.46		
	PLACE1000914	1.05	4.41	3.5	3	6.09	4.22		
20	PLACE1000918	0.54	4.49	1.61	1.82	3.13	1.98		
	PLACE1000927	10.48	12.41	16.9	20.91	23.21	25.47	*	+
	PLACE1000931	0.69	3.44	2.12	2.44	3.94	3.3		
	PLACE1000944	2.55	2.24	4.78	3.84	3.32	2.09		
25	PLACE1000948	0.52	2.31	2.96	2.21	2.72	1.72		
	PLACE1000958	0.12	2.2	1.73	1.11	1.77	2.27		
	PLACE1000972	1.01	3.43	2.89	4.49	5.33	3.75		
	PLACE1000977	2.33	5.67	4.42	2.71	5.33	5.25		
30	PLACE1000979	1.63	8.01	3.93	4.24	5.92	4.57		
	PLACE1000986	3.37	16.51	6.63	6.97	8.75	7.69		
	PLACE1000987	1.76	10.13	4.79	4.17	4.74	5.11		
35	PLACE1001000	4.85	4.62	7.76	6.02	4.25	3.02		
	PLACE1001007	7	6.94	14.66	5.39	3.76	3.47		
	PLACE1001010	0.61	2.04	2.45	2.56	2.73	2.84		
	PLACE1001015	0.88	2.55	1.84	2.36	1.72	2.42		
40	PLACE1001016	1.79	4.54	4.29	6.37	9	6.57	*	+
	PLACE1001022	0.68	6.5	2.45	1.9	2.39	1.29		
	PLACE1001024	1.05	8.89	1.83	1.34	2.49	2.35		
45	PLACE1001036	2.63	10.55	5.42	3.62	5.49	5.43		
	PLACE1001038	50.16	49.81	118.83	82.67	64.83	52.8		
	PLACE1001048	1.07	1.82	0.92	2.39	2.09	1.21		
	PLACE1001054	9.95	10.74	63.88	62.96	79.44	66.71		
50	PLACE1001062	1.45	4.19	3.71	3.12	4.17	3.05		
	PLACE1001063	1.35	3.74	2.2	4.06	3.65	2.74		
	PLACE1001076	0.46	6.48	1	1.86	2.18	1.62		
	PLACE1001081	1.53	7.95	3.33	3.65	5.24	4.8		
55	PLACE1001088	1.32	5.24	1.22	1.42	2.81	1.37		

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	PLACE1000540	5.18	3.93	7.84	5.44	6.9	5.57		
	PLACE1000541	13.59	15.07	48.84	60.62	81.24	41.96		
5	PLACE1000546	0.86	3.61	2.82	4.72	4.63	2.5		
	PLACE1000547	2.16	4.61	3.83	6.31	5.64	5.92	*	+
	PLACE1000560	2.08	5.97	2.1	1.62	2.8	1.72		
	PLACE1000562	2.8	6.23	6.04	8.86	11.26	8.61	*	+
10	PLACE1000564	1.54	6.4	3.07	3.16	4.41	3.43		
	PLACE1000583	3.75	3.28	6.32	6.78	11.53	6.8		
	PLACE1000587	8.52	9.32	12.99	13.64	14.69	9.43		
15	PLACE1000588	1.92	4.36	3.99	8.79	8.15	4.48		
	PLACE1000596	1.99	5.34	4.39	7.8	6.74	4.51		
	PLACE1000599	2.39	5.51	7.05	7.92	7.79	6.46		
	PLACE1000605	5.12	11.43	7.06	14.2	15.1	12.67	*	+
20	PLACE1000610	2.01	6.08	3.54	5.26	4.48	2.94		
	PLACE1000611	13.18	19.13	24.68	16.45	20.37	22.79		
	PLACE1000626	3.19	4.04	8.04	5.71	9.93	8.12		
25	PLACE1000633	1.32	2.12	3.95	4.8	6.06	3.59		
	PLACE1000636	1.15	2.54	1.43	2.72	4.25	2.94		
	PLACE1000653	5.07	8.56	9.29	11.07	11.87	14.08	*	+
	PLACE1000656	4.2	12.9	25.22	16.66	16.71	12.92		
30	PLACE1000663	2	6.43	3.59	2.39	6.61	4.03		
	PLACE1000706	2	6.3	5.04	5.37	7.27	6.26		
	PLACE1000712	3.9	9.52	10.82	10.49	10.07	9.11		
	PLACE1000716	0.98	2.75	3.44	2.44	3.2	2.82		
35	PLACE1000740	2.74	5.28	6.24	8.83	8.13	9.69	*	+
	PLACE1000748	3.35	3.51	6.81	3.12	5.02	4.23		
	PLACE1000749	3.49	6.35	5.94	4.61	4.65	6.02		
40	PLACE1000751	2.71	5.34	4.07	7.81	8.32	8.36	**	+
	PLACE1000755	1.39	6.14	1.93	2.55	5.1	2.96		
	PLACE1000769	2.29	6.8	3.45	3.33	4.58	2.6		
	PLACE1000778	0.87	1.48	1.99	2.05	2.94	2.38		
45	PLACE1000785	9.56	12.21	27.18	28	24.34	29.54		
	PLACE1000786	2.68	4.22	3.63	3.09	3.77	3.7		
	PLACE1000793	4.05	7.21	6.7	6.06	7.6	9.1		
50	PLACE1000795	2.15	5.5	3.99	4.44	5.29	4.31		
	PLACE1000798	0.88	8.44	3.24	3.13	3.8	3.72		
	PLACE1000812	2.13	5.08	4.46	5.06	5.16	6.03		
	PLACE1000823	1.71	5.2	4.89	5.67	7.28	4.84		
55	PLACE1000825	1.6	2.86	2.02	3.77	3.96	3.76	*	+

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	PLACE1000236	0.66	3.33	2.43	2.8	3.63	2.37		
	PLACE1000245	2.92	5.88	6.34	9.34	11.24	10.55	*	+
5	PLACE1000246	5.15	8.27	9.95	3.29	2.55	2.86	*	-
	PLACE1000258	5.4	12.61	13.52	14.88	16.7	14.95		
	PLACE1000288	1.68	7.22	2.96	2.83	4.02	2.48		
	PLACE1000292	3.72	9.02	8.85	10.23	20.58	9.21		
10	PLACE1000302	0.56	1.01	1.39	1.07	0.92	0.62		
	PLACE1000304	1.13	3.26	3.17	3.75	2.32	3.05		
	PLACE1000308	2.54	4.35	4.17	4.42	3.87	1.34		
15	PLACE1000309	2.29	4.02	4	6.72	5.23	7.88	*	+
	PLACE1000312	1.33	3.44	2.48	2.74	3.99	2.5		
	PLACE1000330	0.46	5.76	3.02	1.32	1.93	1.35		
	PLACE1000332	1.02	8.82	2.01	3.01	3.78	1.68		
20	PLACE1000347	2.3	9.48	3.89	2.59	5.81	3.22		-
	PLACE1000351	1.2	1.5	2.87	2.2	2.4	2		
	PLACE1000374	2.01	3.03	7.02	8.89	6.55	6.85		
25	PLACE1000380	2.39	4.27	3.95	4.9	2.12	2.38		
	PLACE1000383	1.03	2.62	1.9	2.53	3.64	2.4		
	PLACE1000397	0.63	4.06	1.89	2.82	3.34	3.47		
	PLACE1000401	1.22	6.39	2.24	2.23	3.05	2.36		
30	PLACE1000406	1.08	8.76	3.4	3.72	4.08	3.64		
	PLACE1000412	1.61	6.38	1.56	1.62	3.45	1.46		
	PLACE1000420	2.59	3.51	4.6	8.95	7.28	5.6	*	+
35	PLACE1000421	0.99	1.3	2.32	2.97	2	1.7		
	PLACE1000423	16.6	23.29	32.85	10.67	8.02	8.09	*	-
	PLACE1000424	1.36	3.09	2.12	3.35	2.65	1.97		
	PLACE1000430	0.77	4.36	1.95	3.51	2.94	3.77		
40	PLACE1000433	1.06	5.9	1.65	1.89	2.6	1.8		
	PLACE1000435	1.39	7.21	4.77	6.22	6.29	4.22		
	PLACE1000437	6.06	10.65	10.14	17.29	20.07	18.79	**	+
	PLACE1000442	3.75	3.85	6.27	7.81	5.7	6.7		
45	PLACE1000444	2.14	3.94	8.96	11.14	11.55	9.8		
	PLACE1000453	5.57	11.03	14.16	10.42	7.4	2.99		
	PLACE1000456	1.25	2.21	1.97	1.33	2.18	1.07		
50	PLACE1000465	2.09	5.63	6.62	12.97	11.8	10.69	**	+
	PLACE1000481	2.32	8.1	3.73	6.89	6.64	6.45		
	PLACE1000492	1.15	4.45	2.95	3.27	3.06	2.81		
	PLACE1000508	1.36	4.64	4	3.91	4.24	3.71		
55	PLACE1000512	4.91	4.29	6.39	8.12	7.8	4.6		

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	OVARC1002156	2.42	3.87	4.19	4.43	3.9	3.65		
	OVARC1002158	0.88	2.63	1.6	2.36	1.57	1.51		
5	OVARC1002165	4.85	6.3	-0.83	10.73	14.03	10.87		
	OVARC1002176	0.86	5.08	3.59	2.01	3.46	2.64		
	OVARC1002178	0.83	5.35	3.12	3.8	5.02	4.25		
	OVARC1002182	1.29	2.89	3.77	2.45	4.64	3.12		
10	OVARC1002185	11.45	13.19	62.79	43.91	53.43	55.56		
	PLACE1000004	1.42	3.23	2.35	3.87	4.25	4.05	*	+
	PLACE1000005	1.18	3.06	3.3	5.27	5.31	4.83	*	+
15	PLACE1000006	2.01	8.33	3.23	4.2	5.44	4.67		
	PLACE1000007	0.97	5.13	2.89	3.03	2.47	2.56		
	PLACE1000014	2.9	8.06	6.26	6.67	8.18	6.55		
	PLACE1000031	0.88	4.81	0.45	2.61	2.71	2.79		
20	PLACE1000033	1.23	2.15	2.75	2.42	3.17	2.56		
	PLACE1000040	3.08	4.43	6.18	7.11	5.54	7.37		
	PLACE1000048	1.83	3.24	2.14	3.32	3.96	3.74		
25	PLACE1000050	2.12	5.36	9.1	9	6.55	8.25		
	PLACE1000061	138.29	147.36	249.77	165.55	233.98	230.37		
	PLACE1000066	14.23	15	19.46	15.86	15.62	18.52		
	PLACE1000075	3.03	6.24	9.08	4.98	6.93	7.11		
30	PLACE1000078	2.1	5.75	5	6.07	6.93	5.19		
	PLACE1000081	1.08	1.88	1.52	1.13	1.89	1.27		
	PLACE1000086	4.97	6.55	11.25	8.1	9.16	7.75		
35	PLACE1000094	0.7	4.18	1.72	1	3.44	2.96		
	PLACE1000101	4.67	8.44	7.7	11.69	10.38	13.65	*	+
	PLACE1000121	0.87	6.29	2.02	1.95	2.85	2.39		
	PLACE1000133	6.65	11.93	17.66	15.19	17.59	21.71		
40	PLACE1000142	1.79	6.03	5.66	2.64	4.77	4.24		
	PLACE1000146	1.95	4.51	2.89	3.71	5.02	2.82		
	PLACE1000163	4.52	5.99	10.71	16.27	10.95	13.78		
45	PLACE1000172	1.12	2.63	1.48	1.81	3.21	2.29		
	PLACE1000181	1.06	3	2.98	2.63	3.75	2.86		
	PLACE1000184	1.17	3.08	1.48	1.87	3.28	2.27		
	PLACE1000185	2.99	6.52	8.47	9.53	9.99	12.03		
50	PLACE1000198	0.78	4.09	2.49	2.19	2.17	1.93		
	PLACE1000213	3.3	5.87	7.36	4.35	5.38	8.09		
	PLACE1000214	1.37	4.29	4.54	5.22	6.72	4.22		
	PLACE1000220	9.61	7.84	16.78	7.48	5.77	4.1		
55	PLACE1000231	2.48	4.1	4.92	4.57	4.65	3.91		

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	OVARC1001873	3.09	3.78	4.68	5.47	4.42	4.73		
	OVARC1001879	1.44	6.08	3.48	2.35	2.93	2.22		
5	OVARC1001880	0.91	7.84	-2.94	3.63	5.78	3.8		
	OVARC1001883	0.99	7.61	3.12	2.61	3.42	2.52		
	OVARC1001900	1.11	7.07	4.03	2.57	3.61	2.89		
	OVARC1001901	0.54	1.84	1.21	2.42	1.37	2.43		
10	OVARC1001911	0.59	1.57	1.66	1.51	1.39	1.55		
	OVARC1001916	1.86	3.13	3.35	4.16	4.61	3.51		
	OVARC1001928	1.45	3.53	1.55	1.58	1.62	1.72		
15	OVARC1001937	5.12	11.69	8.13	17.41	11.63	15.16		
	OVARC1001940	1.1	4.51	3.13	3.72	3.14	2.78		
	OVARC1001942	3.85	7.4	8.03	11.47	13.91	12.77	*	+
	OVARC1001943	7.16	10.07	11.08	9.62	13.85	11.87		
20	OVARC1001949	1.69	3.34	4.15	5.35	4.01	5.55		
	OVARC1001950	1.53	2.41	3.79	6.3	4.35	3.98		
	OVARC1001952	11.3	11.38	53.57	52.33	78.84	38.05		
25	OVARC1001954	1.12	2.99	2.2	3.09	2.67	2.05		
	OVARC1001963	1	4.91	2.89	4.5	3.39	3.21		
	OVARC1001983	3.62	14.16	14.25	20.96	19.21	21.67		
	OVARC1001987	3.12	6.54	5.94	6.08	8.39	8.02		
30	OVARC1001989	1.41	5.2	4.96	4.54	5.59	5.26		
	OVARC1001991	1.74	3.27	4.08	4.57	3.86	3.27		
	OVARC1002005	4.14	3.55	7.66	10.01	9.06	8.2	*	+
	OVARC1002044	3.73	3.94	6.17	6.57	8.32	6.99	*	+
35	OVARC1002046	10.28	16.21	20.07	29.4	37.78	37.02	**	+
	OVARC1002050	1.7	5.6	2.43	3.96	3.82	2.53		
	OVARC1002058	4.23	6.11	4.02	4.69	5.55	5.43		
40	OVARC1002066	11.47	13.5	25.49	26.02	28.69	22.63		
	OVARC1002082	3.6	8.55	8.81	9.6	8.89	6.49		
	OVARC1002091	3.17	5.67	8.37	9.49	5.49	7.64		
	OVARC1002092	1.38	2.72	2.2	4	3.97	1.88		
45	OVARC1002093	1.79	3.1	4.51	5.01	4.44	3.88		
	OVARC1002094	1.55	6.24	4.17	36.42	27.25	28.35	**	+
	OVARC1002107	1.42	4.63	2.69	4.86	5.48	3.6		
50	OVARC1002112	6.17	11.59	8.5	13.47	17.48	11.92		
	OVARC1002126	2.66	6.35	6.68	7.95	6.44	8.79		
	OVARC1002127	0.73	5.04	1.86	1.92	2.61	1.52		
	OVARC1002138	1.4	1.79	1.86	3.16	4.82	2.75	*	+
55	OVARC1002143	0.73	1.51	1.55	1.29	3.03	2.09		

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		OVARC1001611	0.13	5.11	1.24	1.48	2.89	1.79		
		OVARC1001615	0.58	5.42	1.93	1.54	2.56	1.74		
5		OVARC1001636	1.09	3.75	-1.05	2.39	2.05	2.15		
		OVARC1001668	3.77	6.75	10.04	10.5	11.4	10.48		
		OVARC1001702	1.18	2.21	2.42	3.86	2.07	2.25		
10		OVARC1001703	2.82	3.18	2.97	2.64	4.71	4.65		
		OVARC1001710	3.58	7.03	8.67	8.01	6.28	9.55		
		OVARC1001711	1.96	7.3	3.36	5.01	4.15	5.4		
		OVARC1001713	9.17	11.54	44.65	36.47	51.48	43.79		
15		OVARC1001725	1.01	5.45	6.11	2.56	3.77	4.01		
		OVARC1001726	1.64	4.48	3.23	4.97	5.6	5.18		
		OVARC1001727	1.4	2.41	1.52	1.43	2.35	1.14		
20		OVARC1001731	120.62	110.86	255.43	140.73	139.03	74.2		
		OVARC1001735	1.29	3.44	3.54	3.75	3.25	2.89		
		OVARC1001741	3.3	4.73	15.28	13.09	12.93	13.17		
		OVARC1001745	2.72	5.39	6.83	9.17	10.23	8.89	*	+
25		OVARC1001759	3.31	9.01	6.31	7.61	7.61	11.95		
		OVARC1001762	3.96	7.78	6.38	10.3	11.01	13.4	*	+
		OVARC1001766	5.33	7.8	11.99	15.56	16.86	15.33	*	+
		OVARC1001767	0.94	3.76	1.18	1.97	1.96	2.41		
30		OVARC1001768	3.31	3.86	3.76	5.35	4.59	3.54		
		OVARC1001770	3.04	6.58	9.98	11.46	9.08	9.46		
		OVARC1001776	2.11	4.7	3.5	4.72	3.64	2.92		
35		OVARC1001791	1.13	4.77	3.54	3.07	3.12	3.42		
		OVARC1001795	0.89	6.19	1.24	2.31	2.87	2.11		
		OVARC1001798	2.81	12.11	7.57	9.72	11.93	9.04		
		OVARC1001802	1.73	11.64	4.9	5.6	5.93	4.01		
40		OVARC1001805	1.92	6.96	2.58	3.62	4.59	3.51		
		OVARC1001807	1.9	2.53	4.18	3.06	3.12	2.46		
		OVARC1001809	12.38	14.06	76.32	55.87	81.41	52.83		
45		OVARC1001812	1.44	3.39	3.15	3.23	4.63	3.71		
		OVARC1001813	1.61	4.29	2.33	2.93	3.98	2.51		
		OVARC1001820	1.67	7.15	3.21	3.47	3.76	3.22		
		OVARC1001828	0.78	6.85	2.36	1.91	3.23	2.2		
50		OVARC1001833	1.07	8.12	2.02	2.4	2.1	1.92		
		OVARC1001839	1.56	8.43	2.98	2.91	3.59	1.15		
		OVARC1001846	1.91	1.38	2.9	2.15	2.11	1.8		
		OVARC1001849	1.21	2.52	2.42	5.79	3.69	4.03	*	+
55		OVARC1001861	1.46	3.56	2.73	2.78	2.5	2.09		

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	OVARC1001372	0.96	4.23	3.33	1.52	2.77	1.95		
	OVARC1001376	1.82	5.1	3.62	5.79	5.79	4.18		
5	OVARC1001381	4.51	6.44	9.94	10.95	12.91	11.21		
	OVARC1001391	0.5	1.62	1.44	1.88	1.27	1.26		
	OVARC1001392	2.12	4.69	6.14	11.96	12.7	7.79	*	+
	OVARC1001399	0.98	3.59	2.16	1.77	2.54	1.1		
10	OVARC1001417	1.01	3.07	1.76	2.39	3.61	2.81		
	OVARC1001419	2.47	5.4	3.06	4.39	4.03	3.16		
	OVARC1001425	2.29	5.58	5.15	8.76	8.5	8.07	*	+
15	OVARC1001436	1.37	5.85	2.54	2.9	3.57	3.18		
	OVARC1001442	0.64	4.84	1.39	2.27	1.52	0.69		
	OVARC1001451	3.09	2	3.89	5.18	5.98	4.75	*	+
	OVARC1001452	1.35	2.41	2.87	2.96	4.69	3.13		
20	OVARC1001453	1.21	2.84	1.88	2.3	1.82	1.57		
	OVARC1001476	10.67	14.38	16.52	17.22	12.9	13.09		
	OVARC1001480	0.93	4.73	1.5	3.1	2.98	2.21		
25	OVARC1001489	0.97	6.89	2.51	3.01	2.83	2.09		
	OVARC1001493	2.09	6.59	3.75	7.38	8.78	10.48	*	+
	OVARC1001496	4.65	9.58	8.63	10.74	7.37	11.03		
	OVARC1001499	1.24	1.18	2.6	3.47	2.68	2.84		
30	OVARC1001506	2.9	2.7	5.31	5.33	6.73	5.48		
	OVARC1001509	1.73	3.89	3.69	4.07	4.59	3.07		
	OVARC1001510	0.16	3.69	1.42	1.94	1.73	0.86		
35	OVARC1001516	2.57	5.78	3.85	6.04	4.97	4.39		
	OVARC1001525	0.53	4.76	2.12	1.94	2.01	1.81		
	OVARC1001542	8.8	12.86	13.01	15.91	13.63	17.23		
	OVARC1001544	2.14	6.6	6.72	7.54	8.33	6.22		
40	OVARC1001546	4.08	4.32	4.6	6.12	5.31	7.23	*	+
	OVARC1001547	1.29	2.53	1.68	2.44	1.85	2.22		
	OVARC1001555	10.39	16.51	68.77	48.66	65.39	56.39		
45	OVARC1001560	3.35	4.91	5.52	5.36	4.93	5.34		
	OVARC1001569	1.63	4.75	4.79	5.92	5.19	5.1		
	OVARC1001570	3.96	7.9	6.93	7.72	10.7	8.55		
	OVARC1001577	1.68	5.89	5.41	8.61	6.9	10.2		
50	OVARC1001578	0.25	3.47	-0.19	-0.47	0.24	0.15		
	OVARC1001596	12.13	11.65	14.23	13.51	14.82	27.15		
	OVARC1001600	1.13	2.9	1.48	2.81	2.67	3.67		
	OVARC1001607	6.22	7.72	10.91	13.42	14.01	13.45	*	+
55	OVARC1001610	1.81	5.25	2.84	4.25	2.66	2.7		

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	OVARC1001173	2.09	5.08	5.1	4.32	6.75	5.49		
	OVARC1001176	22.57	21.48	89.96	76.74	102.3	70.7		
5	OVARC1001180	3.01	4.58	12.7	11.81	10.77	9.56		
	OVARC1001188	2.66	3.7	3.95	3.62	3.44	4.16		
	OVARC1001200	1.52	4.56	3.62	3.47	2.9	2.96		
10	OVARC1001202	3.75	6.65	6.53	9.26	7.79	10.23	*	+
	OVARC1001206	1.52	5.52	1.15	1.59	1.13	1.9		
	OVARC1001209	4.89	8.92	27.46	24.69	30.38	24.27		
	OVARC1001219	1.81	6.36	4.71	5.33	3.95	3.62		
15	OVARC1001222	2.5	8.36	5.01	3.2	4.34	5.63		
	OVARC1001232	2.91	4.18	7.74	6.75	6.02	5.65		
	OVARC1001240	2.05	3.27	6.84	5.55	5.06	5.4		
20	OVARC1001243	0.94	2.59	1.76	3.64	2.64	1.86		
	OVARC1001244	9.07	12.05	18	21.61	18.57	26.62		
	OVARC1001246	30.48	50.95	48.51	80.54	100.83	101.88	**	+
	OVARC1001247	3.64	9.86	7.7	6.57	7.02	4.49		
25	OVARC1001260	1.05	9.07	1.85	2.62	2.65	1.85		
	OVARC1001261	4.23	10.5	6.99	3.46	2.08	2.94		
	OVARC1001268	24.4	19.69	52.37	32.58	35.32	14.16		
30	OVARC1001270	14.46	15.1	20.83	9.69	9.8	8.65	*	-
	OVARC1001271	2.62	3.62	3.88	3.95	7.02	4.26		
	OVARC1001282	0.88	3.02	3.09	1.37	1.59	2		
	OVARC1001296	3.02	8.06	2.3	3.04	4.11	5.41		
35	OVARC1001306	1.48	8.27	2.4	2.04	2.29	3.82		
	OVARC1001314	0.49	8.47	1.57	1.06	1.79	1.32		
	OVARC1001316	2.77	7.17	4.81	5.48	8.11	5.36		
	OVARC1001329	6.12	6.18	21.11	17.09	19.29	16.22		
40	OVARC1001330	0.2	1.89	1.38	1.22	1.35	1.42		
	OVARC1001336	1.92	3.7	3.59	5.67	4.09	4.02		
	OVARC1001338	0.26	2.87	0.86	2.49	1.71	1.07		
45	OVARC1001339	12.07	18.29	22.73	33.65	32.72	37.29	**	+
	OVARC1001340	0.72	4.83	1.23	1.33	2.44	1.3		
	OVARC1001341	4.35	9.25	6.77	7.94	11.38	9.69		
	OVARC1001342	90.37	98.53	136.12	129.68	163.22	127.78		
50	OVARC1001344	2.1	2.51	6.27	6.52	6.89	6.2		
	OVARC1001357	5.61	8.93	16.02	15.52	11.34	11.69		
	OVARC1001359	8.96	12.4	16.15	21.66	13.84	10.6		
	OVARC1001360	0.44	2.52	0.99	1.97	2.6	1.62		
55	OVARC1001369	1.56	5.66	1.89	3.41	1.88	3.1		

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		OVARC1001032	1.42	5.34	1.89	1.83	2.82	1.92		
		OVARC1001034	4.44	5.58	6.51	3.29	5.21	3.77		
5		OVARC1001038	3.62	5.03	7.4	10.3	10.88	8.61	*	+
		OVARC1001040	2.63	3.77	6.93	5.25	6.51	4.25		
		OVARC1001041	4.54	8.03	12.87	8.57	12.25	9.4		
10		OVARC1001044	1.05	2.92	1.83	1.96	2.43	1.84		
		OVARC1001049	3.78	8.78	10.67	10.65	11.87	10.26		
		OVARC1001051	40.95	55.97	80.66	66.89	109.71	87.49		
		OVARC1001054	1.22	4.06	3.22	2.86	4.19	1.93		
15		OVARC1001055	2.13	3.38	3.82	4.32	5.61	5.22	*	+
		OVARC1001062	5.8	6.15	12.54	8.04	9.94	9.57		
		OVARC1001065	8.85	13.63	51.33	51.41	60.3	56.97		
20		OVARC1001068	2.82	5.62	4.76	4.72	4.02	5.52		
		OVARC1001072	0.73	4.18	4.41	3.2	3.71	3.07		
		OVARC1001073	0.92	5.7	2.65	2.91	2.54	1.79		
		OVARC1001074	0.81	4.66	3.31	1.87	2.95	2.04		
25		OVARC1001078	2	5.12	2.79	3.57	3.08	2.83		
		OVARC1001085	2.41	2.83	3.66	5.54	5.02	6.36	**	+
		OVARC1001086	1.97	3.17	2.85	3.98	2.83	4.13		
30		OVARC1001091	16.24	19.32	92.73	76.48	96.74	77.99		
		OVARC1001092	4.62	5.35	7.22	9.69	7.84	6.05		
		OVARC1001104	1.05	4.37	2.66	3.16	2.58	2.03		
		OVARC1001107	11.59	15.6	40.28	31.21	49.49	42.22		
35		OVARC1001113	1.04	5.81	1.59	2.46	3.05	2.39		
		OVARC1001117	2.71	6.63	4.31	4.67	5.74	2.67		
		OVARC1001118	2.38	3.69	7.08	7.36	6.91	8.28		
		OVARC1001125	2.02	2.9	3	5.92	4.97	5.9	**	+
40		OVARC1001129	2.61	4.58	3.19	8.46	9.43	11	**	+
		OVARC1001132	1.7	6.48	2.66	3.69	4.26	4.66		
		OVARC1001138	9.52	15.82	23.8	48.95	45.16	44.97	**	+
45		OVARC1001141	1.68	4.97	3.48	3.77	3.68	3.84		
		OVARC1001154	18.31	29.49	68.39	60.43	83.49	65.64		
		OVARC1001161	2.49	5.55	6.15	7.03	6.69	5.89		
		OVARC1001162	2.2	3.13	5.34	5.09	4.86	5.26		
50		OVARC1001163	0.69	3.59	2.77	2.2	3.98	2.54		
		OVARC1001167	3.03	4.57	7.69	10.19	12.95	9.3	*	+
		OVARC1001169	0.74	4.87	2.68	2.47	1.91	2.06		
55		OVARC1001170	7.81	15.04	17.59	14.61	19.45	14.99		
		OVARC1001171	15.57	17.71	24.31	16.12	23.34	22.51		

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	OVARC1000622	3.86	4.59	11.57	12.7	11.13	11.6		
	OVARC1000636	1.64	3.79	4.58	4.19	4.03	5.09		
5	OVARC1000640	1.97	4.72	3.93	4.21	3.67	3.09		
	OVARC1000649	9.69	14.8	53.54	53.32	64.51	52.67		
	OVARC1000661	1.14	9.33	2.99	5.34	5.24	5.24		
	OVARC1000677	8.53	10.16	14.87	11.77	10.98	15.47		
10	OVARC1000678	1.21	4.49	2.71	3.28	4.17	3.26		
	OVARC1000679	2.86	3.25	4.09	5.29	5.16	6.25	*	+
	OVARC1000681	1.47	1.55	3.2	2.41	2.22	1.71		
15	OVARC1000682	10.09	11.33	50.91	33.79	47.49	44.31		
	OVARC1000689	3.81	7.1	19.94	20.18	22.12	21.83		
	OVARC1000700	1.8	10.37	3.18	4.98	4.37	5.14		
	OVARC1000703	1.74	7.18	5.35	6.4	6.64	7.77		
20	OVARC1000722	10.59	11.92	47.93	43.41	60.06	39.34		
	OVARC1000726	1.44	3.48	4.62	4.88	5.89	3.58		
	OVARC1000727	1.93	2.09	4.13	3.78	3.79	3.89		
25	OVARC1000730	5.95	5.86	9.01	4.07	4.16	5.62		
	OVARC1000741	4.85	6.13	8.74	15.19	10.58	13.71	*	+
	OVARC1000746	0.89	3.61	2.43	2.06	2.9	2.84		
	OVARC1000764	1.76	4.93	4.77	5.35	7.01	5.44		
30	OVARC1000769	1.13	4.3	3.6	3.76	4.42	5.2		
	OVARC1000771	2.42	6.28	2.3	4.02	4.81	3.71		
	OVARC1000773	19.09	24.7	31.93	44.69	56.24	46.24	*	+
35	OVARC1000775	11.67	8.94	16.44	12.16	8.7	4.26		
	OVARC1000778	2.37	3.89	5.69	4.59	6.23	4.92		
	OVARC1000779	0.8	2.02	1.85	2.23	2.45	1.46		
	OVARC1000781	1.67	5.05	4.16	6.37	3.45	5.07		
40	OVARC1000787	1.64	4.79	4.22	2.97	5.44	3.25		
	OVARC1000789	7.62	14.23	16.39	24.95	29.69	25.94	**	+
	OVARC1000800	2.91	10.72	5.72	6.41	10.65	6.2		
45	OVARC1000802	1.55	8.77	2.97	2.99	5.34	2.76		
	OVARC1000810	3.37	3.54	8.29	6.66	7.99	7.21		
	OVARC1000811	2.41	2.73	7.5	4.88	3.67	4.95		
	OVARC1000814	3.44	4.55	9.03	7.92	11.3	9.05		
50	OVARC1000816	7.64	10.41	12.41	10.99	10.58	14.11		
	OVARC1000817	1.18	3.38	1.27	1.71	2.14	1.55		
	OVARC1000834	2.46	8.3	3.39	4.84	5.81	4.01		
	OVARC1000846	2.23	10.02	5.35	7.38	9.66	7.72		
55	OVARC1000850	1.74	8.37	3.38	3.39	2.51	3.1		

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	OVARC1000414	1.31	2.22	3.7	4.5	3.78	3.71		
	OVARC1000420	1.44	2.76	3.29	3.3	2.59	2.1		
5	OVARC1000421	1.42	2.65	3.33	3.96	4.21	4.99	*	+
	OVARC1000427	25.78	27.02	130.06	156.9	215.67	142.19		
	OVARC1000431	10.51	17.6	19.12	33.66	31.78	25.78	*	+
	OVARC1000437	3.14	6.37	7.31	5.97	7.63	6.36		
10	OVARC1000439	5.81	10.95	13.82	21.81	23.01	21.52	**	+
	OVARC1000440	2.56	3.74	5.01	7.47	9.31	7.08	*	+
	OVARC1000442	2.34	2.38	6.81	6.66	9.5	8.12		
15	OVARC1000443	2.09	2.2	2.88	3.29	3.41	2.62		
	OVARC1000461	1.11	2.84	2.2	2.55	1.12	2.14		
	OVARC1000465	3.27	5.01	3.51	3.94	4.62	3.95		
	OVARC1000466	1.94	5.47	5.9	6.54	10.13	6.76		
20	OVARC1000467	1.01	5.08	2.41	3.65	2.98	3.78		
	OVARC1000470	1.13	5.81	3.03	3.18	4.02	3.78		
	OVARC1000473	1.81	1.95	2.65	2.44	4.16	1.39		
25	OVARC1000479	5.67	5.88	9.88	10.35	14.26	6.88		
	OVARC1000484	3.99	5.74	6.54	8.66	9.93	6.87		
	OVARC1000486	3.17	4.71	4.49	5.74	4.93	4.28		
	OVARC1000496	0.93	3.55	0.66	0.31	1.07	0.62		
30	OVARC1000520	0.84	5.89	1.18	1.32	2.27	2		
	OVARC1000522	4.1	7.19	12	13.85	14.03	10.34		
	OVARC1000526	1.96	7.04	3.75	5.93	5.48	4.69		
	OVARC1000529	2.38	2.57	4.44	3.66	4.16	3.08		
35	OVARC1000533	3.3	4.66	7.95	8.89	12.29	7.83		
	OVARC1000543	0.84	2.44	2.06	2.57	3.08	2.72		
	OVARC1000550	0.75	3.68	2.32	2.82	4.04	2.34		
40	OVARC1000553	2.1	5.63	7.02	7.72	7.69	6.95		
	OVARC1000556	5.77	15.21	11.77	8.95	13.91	8.96		
	OVARC1000557	0.83	5.12	1.4	1.61	2.29	1.88		
	OVARC1000561	3.48	7.38	9.26	13	17.66	15.09	*	+
45	OVARC1000564	8.89	9.02	10.44	17.84	11.31	16.69	*	+
	OVARC1000573	1.87	3.68	4.86	5.75	5.32	3.83		
	OVARC1000576	24.12	29.23	124.94	83.09	93.83	98.58		
50	OVARC1000578	2.43	4.6	5.53	8.64	4.46	3.93		
	OVARC1000581	0.34	3.28	1.15	1.75	1.27	1.23		
	OVARC1000586	22.54	28.9	41.17	34.58	43.39	40.93		
	OVARC1000588	0.74	5.23	2.03	2.75	3.72	2.05		
55	OVARC1000605	1.98	2.62	2.88	4.47	4.23	3.87	**	+

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		OVARC1000191	1. 01	3. 8	2. 63	3. 12	2. 85	2. 54		
		OVARC1000198	2. 09	3. 59	4. 32	5. 62	5. 12	5. 06	*	+
5		OVARC1000208	6. 49	10. 37	22. 5	17. 79	24. 54	22. 02		
		OVARC1000209	7. 99	13. 69	22. 82	23. 42	27. 81	29. 16		
		OVARC1000212	2. 47	5. 63	3. 59	4. 76	5. 03	4. 88		
10		OVARC1000216	1. 72	4. 96	4. 36	15. 43	11. 3	12. 54	**	+
		OVARC1000240	2. 98	3. 53	8. 13	5. 39	5. 46	4. 87		
		OVARC1000241	1. 29	2. 47	3. 18	2. 65	3. 17	1. 4		
		OVARC1000249	4. 14	5. 43	8. 17	5. 46	5	6. 13		
15		OVARC1000254	33. 15	39. 39	100. 99	100. 41	131. 42	100. 89		
		OVARC1000255	0. 85	4. 83	2. 51	2. 98	2. 45	1. 95		
		OVARC1000267	2. 37	6. 41	6. 71	6. 66	7. 16	7. 31		
20		OVARC1000275	79. 02	93. 7	161. 08	199. 43	240. 76	175. 96	*	+
		OVARC1000287	226. 67	224. 66	236. 08	433. 91	512. 76	470. 31	**	+
		OVARC1000288	3. 2	4. 25	7. 38	6. 23	5. 32	4. 47		
		OVARC1000298	8. 96	10. 09	19. 62	13. 37	7. 19	9. 6		
25		OVARC1000302	1. 12	2. 14	2. 13	2. 47	1. 85	2		
		OVARC1000304	1. 09	2. 68	3. 23	5. 02	3. 41	6		
		OVARC1000307	2. 95	6. 19	4. 74	7. 59	4. 7	6. 29		
		OVARC1000309	1. 18	7. 16	3. 22	3. 24	2. 85	2. 4		
30		OVARC1000312	2. 83	11. 64	6. 03	4. 17	5. 4	2. 46		
		OVARC1000313	10. 48	19. 25	14. 81	9. 39	17. 54	22. 17		
		OVARC1000321	31. 6	24. 05	47. 79	30. 5	31. 37	15. 43		
35		OVARC1000326	1. 52	2. 3	3. 9	3. 84	3. 17	2. 79		
		OVARC1000327	1. 52	3. 28	4. 24	3. 13	1. 49	2. 46		
		OVARC1000331	2. 22	4. 72	2. 41	4. 33	4. 45	4. 58		
		OVARC1000335	2. 3	5. 84	4. 02	2. 72	5. 16	4. 75		
40		OVARC1000347	1. 83	8. 18	6. 24	7. 35	9. 24	8. 44		
		OVARC1000348	1. 61	10. 62	3. 73	2. 84	4. 59	3. 05		
		OVARC1000363	3. 7	9. 61	6. 51	7. 31	11. 52	6. 83		
45		OVARC1000377	1. 07	2. 09	2. 43	2. 28	2. 51	2. 45		
		OVARC1000382	3. 34	3. 39	4. 33	5. 07	2. 52	1. 03		
		OVARC1000384	4. 2	5. 42	8. 35	5. 32	4. 4	6. 04		
		OVARC1000401	0. 62	3. 63	2. 09	3. 35	4. 08	3. 64		
50		OVARC1000406	18. 98	23. 3	49. 12	57. 09	74. 48	54. 63	*	+
		OVARC1000407	1. 99	6. 28	3. 99	4. 11	6. 42	3. 16		
		OVARC1000408	27. 5	38. 45	70. 39	74. 84	111. 17	71. 25		
55		OVARC1000410	6. 83	12. 72	10. 41	4. 78	6. 65	5. 01		
		OVARC1000411	0. 91	1. 5	2. 6	3. 49	3	2. 22		

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	OVARC1000006	3.57	3.57	6.58	9.04	7.88	4.73		
	OVARC1000013	6.51	6.51	9.19	7.32	8.36	8.33		
5	OVARC1000014	3.39	3.39	5.02	4.23	5.02	4.17		
	OVARC1000017	3.11	3.11	6.81	4.2	4.45	2.72		
	OVARC1000026	24.79	24.79	32.1	56.82	69.34	44.53	*	+
	OVARC1000035	11.11	11.11	20.26	20.41	23.65	15.36		
10	OVARC1000037	8.73	8.73	19.12	15.64	9.13	15.9		
	OVARC1000058	6.06	6.06	11.69	13.84	7.56	10.6		
	OVARC1000060	1.89	1.89	6.28	5.98	5.24	5.13		
15	OVARC1000068	2.38	2.38	5.33	5.31	3.56	2.42		
	OVARC1000069	74.66	74.66	101.53	75.95	84.36	86.42		
	OVARC1000071	4.4	4.4	4.77	6.47	5.35	4.04		
	OVARC1000075	55.43	55.43	125.63	120.89	150.97	117.03		
20	OVARC1000083	9.58	9.58	9.24	13.12	12.7	10.64	*	+
	OVARC1000085	106.6	90.9	156.14	214.2	177.05	273.14	*	+
	OVARC1000086	3.98	6.82	9.23	11.98	11.3	14.09	*	+
25	OVARC1000087	1.51	2.83	1.79	4.03	3.57	3.35	*	+
	OVARC1000090	1.48	4.1	6.14	10.88	9.58	8.79	*	+
	OVARC1000091	4.88	8.33	8.01	7.99	7.76	6.82		
	OVARC1000092	2.83	6.81	4.18	4.68	6.25	4.85		
30	OVARC1000105	9.73	14.86	17.21	26.29	25.62	22.88	*	+
	OVARC1000106	26.02	23.03	46.38	66.36	50.1	53.01	*	+
	OVARC1000109	9.12	13.08	18.04	16.72	12.91	17.46		
35	OVARC1000113	4.12	6.25	6.53	6.83	8.19	7.65		
	OVARC1000114	2.14	3.44	5.77	5.94	5.86	4.98		
	OVARC1000133	2.53	4.96	6.36	4.05	4.97	2.95		
	OVARC1000137	6.14	10.05	13.51	13.3	18.59	14.39		
40	OVARC1000139	14.75	20.77	83.44	71.14	98.1	69.29		
	OVARC1000145	0.72	6.64	2.89	1.78	2.42	2		
	OVARC1000148	5.09	4.98	7.88	4.91	5.32	7.91		
45	OVARC1000151	1.41	2.11	2.4	3.58	4.08	3.58	**	+
	OVARC1000157	10.99	14.16	17.51	21.21	25.06	22.76	*	+
	OVARC1000162	1.22	4.4	2.5	2.93	2.49	2.59		
	OVARC1000168	1.98	8.46	6.2	8.01	9.61	9.96		
50	OVARC1000169	32.03	45.07	49.48	70.63	69.6	89.08	*	+
	OVARC1000178	0.84	5.08	2.53	3.37	3.18	2.78		
	OVARC1000182	0.8	3.3	1.42	2.02	1.95	1.78		
	OVARC1000186	2.51	3.72	3.23	5.95	3.27	4.77		
55	OVARC1000188	1.04	2.67	2.33	2.48	2.87	1.9		

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	NT2RP4002850	5.07	5.07	12.04	14.36	12.63	8.06		
	NT2RP4002874	5.17	5.17	6.67	3.41	5.14	1.96		
5	NT2RP4002884	28.81	28.81	49.75	43.57	74.75	52.87		
	NT2RP4002888	5.55	5.55	4.83	3.67	4.32	3.08	*	-
	NT2RP4002891	5.48	5.48	15.79	13.16	19.42	11.91		
10	NT2RP4002894	12.04	12.04	24.47	18.44	12.76	16.4		
	NT2RP4002896	5.54	5.54	12.2	8.96	6.18	7.78		
	NT2RP4002905	1.71	1.71	4.27	2.32	3.58	1.28		
	NT2RP4002907	5.11	5.11	7.62	6.94	10.72	1.41		
15	NT2RP5003459	68.11	68.11	133.25	154.61	146.15	164.37	*	+
	NT2RP5003461	7.34	7.34	10.14	10.85	14.36	8		
	NT2RP5003471	106.6	106.6	168.71	124.4	148.85	112.14		
20	NT2RP5003477	2.71	2.71	2.62	2.59	2.33	1.9		
	NT2RP5003487	157.44	157.44	424.89	292.71	256.56	354.93		
	NT2RP5003492	3.1	3.1	4.91	5.25	6.17	5.91	*	+
	NT2RP5003500	1.5	1.5	3.28	2.38	2.54	2.59		
25	NT2RP5003506	4.96	4.96	9.3	7.83	10.37	9.04		
	NT2RP5003512	2.21	2.21	4.35	2.63	3.46	2.15		
	NT2RP5003522	4.1	4.1	5.97	4.62	4.19	2.34		
30	NT2RP5003524	4.38	4.38	3.86	1.61	1.54	0.84	**	-
	NT2RP5003527	24.72	24.72	71.27	76.81	87.24	60.59		
	NT2RP5003531	7.16	7.16	17.2	15.58	14.06	14.11		
	NT2RP5003534	2.68	2.68	5.49	5.54	6.82	4.55		
35	NT2RP6000020	8.69	8.69	19.96	14.65	15.13	16.29		
	NT2RP6000022	3.19	3.19	4.05	4.06	3.96	2.44		
	NT2RP6000050	3.95	3.95	3.99	4.98	5.82	2.88		
	NT2RP6000063	3.91	3.91	6.04	3.61	2.52	2.56		
40	NT2RP6000074	5.38	5.38	4.88	3.41	3.27	2.17	**	-
	NT2RP6000083	7.76	7.76	11.18	11.49	16	9.91		
	NT2RP6000100	2.49	2.49	4.58	4.04	4.71	3.3		
45	NT2RP6000123	1.94	1.94	3.29	5.1	4.26	4.22	*	+
	NT2RP6000129	1.9	1.9	4.47	4.06	4.27	2.74		
	NT2RP6000147	3.75	3.75	11.74	10.8	11.03	7.48		
	NT2RP6000163	2.62	2.62	4.23	2.28	1.95	1.71		
50	NT2RP6000181	8.03	8.03	12.4	9.44	13.25	9.01		
	NT2RP6000182	5.44	5.44	6.42	4.82	5.56	3.88		
	OVARC1000001	4.97	4.97	5.24	6.04	7.48	2.35		
55	OVARC1000003	3.21	3.21	8.31	8.51	7.66	7.05		
	OVARC1000004	9.87	9.87	116.19	88.04	109.99	85.44		

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	NT2RP4002288	6.42	6.42	12.57	13.29	14.36	11.97		
	NT2RP4002290	7.55	7.55	7.61	7.96	7.67	5.87		
5	NT2RP4002298	3.92	3.92	4.18	5.54	5.03	4.18		
	NT2RP4002306	2.38	2.38	5.79	2.97	5.77	2.64		
	NT2RP4002308	2.04	2.04	6.03	5.31	5.23	4.1		
	NT2RP4002336	2.71	2.71	6.33	3.71	4.19	4.63		
10	NT2RP4002340	1.09	1.09	3.96	1.28	2.75	0.49		
	NT2RP4002361	2.77	2.77	5.78	3.73	4.03	2.48		
	NT2RP4002367	2.27	2.27	5.84	3.23	2.48	2.77		
15	NT2RP4002368	9.87	9.87	17.2	18.26	19.27	16		
	NT2RP4002377	3.3	3.3	23.8	25.46	30.75	23.93		
	NT2RP4002408	2.22	2.22	3.87	3.75	6.37	4.11		
	NT2RP4002425	2.84	2.84	5.81	8.24	7.98	5.23		
20	NT2RP4002432	12.33	12.33	85.4	61.06	72.53	67.82		
	NT2RP4002447	2.97	2.97	7.68	3.96	5.4	4.59		
	NT2RP4002451	5.48	5.48	6.2	5.84	5.85	6.83		
25	NT2RP4002461	9.8	9.8	32.09	32.76	38.91	29.04		
	NT2RP4002486	3.5	3.5	6.71	2.47	4.15	2.87		
	NT2RP4002517	3.65	3.65	9.11	7.02	8.53	7.18		
	NT2RP4002556	4.29	4.29	3.91	5.68	10.03	6.41		
30	NT2RP4002569	3.36	3.36	7.36	4.93	5.29	3.42		
	NT2RP4002587	2.26	2.26	4.19	2.8	3.4	2.02		
	NT2RP4002591	2.21	2.21	4.89	2.89	4.5	3.08		
35	NT2RP4002607	1.43	1.43	3.34	2.87	4.63	1.58		
	NT2RP4002627	17.83	17.83	61.9	55.9	76.17	65.3		
	NT2RP4002628	7.28	7.28	15.48	14.53	23.95	12.54		
	NT2RP4002630	4.19	4.19	5.25	6.72	9.4	7.16	*	+
40	NT2RP4002639	9.43	9.43	70.25	52.38	77.24	57.28		
	NT2RP4002641	1.58	1.58	9.03	3.94	4.07	4.1		
	NT2RP4002658	114.62	114.62	166.93	76.49	34.96	109.83		
45	NT2RP4002669	3.5	3.5	5.67	5.4	5.33	4.68		
	NT2RP4002677	6.24	6.24	9.41	10.14	7.99	13.62		
	NT2RP4002715	8.42	8.42	34.92	40.1	48.46	32.3		
	NT2RP4002750	2.6	2.6	8.29	1.68	2.04	1.33		
50	NT2RP4002784	3.71	3.71	9.51	9.44	11.22	7.06		
	NT2RP4002791	4.91	4.91	9.44	4.88	9.76	5.33		
	NT2RP4002811	1.63	1.63	6.38	3.17	2.95	3.43		
	NT2RP4002830	4.26	4.26	7.45	3.9	5.9	5.46		
55	NT2RP4002832	2.12	2.12	3.13	2.38	5.59	2.54		

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	NT2RP4002018	6.19	6.19	13.71	10.47	11.39	10.36		
	NT2RP4002035	5.35	5.35	5.95	5.4	4.17	2.54		
5	NT2RP4002043	7.1	7.1	10.8	9.64	12.2	6.5		
	NT2RP4002046	9.74	9.74	20.08	21.94	15.28	17.11		
	NT2RP4002047	8.37	8.37	19.18	22.28	24.07	28.83	*	+
10	NT2RP4002052	5.78	5.78	10.36	9.02	9.36	9.37		
	NT2RP4002056	32.58	32.58	71.49	58.09	76.58	49.75		
	NT2RP4002057	6.37	6.37	11.06	12.13	13.58	7.41		
	NT2RP4002058	3.85	3.85	6.6	4.1	4.2	3.22		
15	NT2RP4002064	5.93	5.93	4.39	2.6	4.16	2.07	*	-
	NT2RP4002071	6.67	6.67	7.07	6.95	10.06	6.27		
	NT2RP4002075	1.16	1.16	2.11	2.27	2.35	1.27		
20	NT2RP4002078	2.25	2.25	8.63	6.86	8.24	4.97		
	NT2RP4002081	8.11	8.11	26.15	18.73	18.78	19.42		
	NT2RP4002083	1.39	1.39	5.25	3.36	3.16	1.88		
	NT2RP4002099	3.26	3.26	4.73	2.84	3.56	2.51		
25	NT2RP4002106	10.35	10.35	20.34	22.36	25.93	20.55		
	NT2RP4002111	11.7	11.7	12.37	19.77	30.44	17.72		
	NT2RP4002112	6.15	6.15	10.97	8.9	8.34	3.22		
30	NT2RP4002116	12.6	12.6	47.19	37.43	41.25	28.65		
	NT2RP4002122	5.34	5.34	9.29	14.84	14.86	12.67	**	+
	NT2RP4002126	6.42	6.42	14.44	16.82	14.35	10.42		
	NT2RP4002133	7.56	7.56	20.82	29.17	26.14	21		
35	NT2RP4002136	3.63	3.63	5.74	4.89	5.38	2.69		
	NT2RP4002139	26.89	26.89	31.12	60.65	61.92	32.88		
	NT2RP4002174	139.27	139.27	232.64	240.71	275.01	193.19		
40	NT2RP4002185	7.77	7.77	13.2	12.36	19.06	11.58		
	NT2RP4002186	4.5	4.5	9.83	7.82	4.72	6.78		
	NT2RP4002187	15.42	15.42	32.13	26.94	19.84	21.17		
	NT2RP4002188	3.01	3.01	8.34	8.3	7.75	6.41		
45	NT2RP4002199	1.85	1.85	3.73	2.6	2.91	3.78		
	NT2RP4002206	2.08	2.08	3.39	2.48	2.34	1.29		
	NT2RP4002210	3.13	3.13	4.75	2.02	2.86	0.98		
	NT2RP4002222	4.2	4.2	6.63	5.56	6.28	4.16		
50	NT2RP4002241	7.97	7.97	8.24	10.82	11.75	7.19		
	NT2RP4002248	5.08	5.08	16.69	10.74	8.9	8.13		
	NT2RP4002250	1.54	1.54	3.22	0.73	1.69	0.56		
55	NT2RP4002259	4.86	4.86	9.82	3.21	4.85	1.75		
	NT2RP4002268	16.62	16.62	29.54	28.9	28.18	25.68		

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	NT2RP4001803	3.35	3.35	3.6	3.94	4.78	3.67		
	NT2RP4001805	2.64	2.64	3.57	3.64	2.47	2.95		
5	NT2RP4001809	4.84	4.84	26.35	18.18	23.17	11.33		
	NT2RP4001817	11.55	11.55	19.09	9.5	10.78	12.71		
	NT2RP4001822	2.09	2.09	5.36	3.73	5.11	3.33		
	NT2RP4001823	1.91	1.91	3.95	1.14	3.34	1.24		
10	NT2RP4001827	14.88	14.88	25.96	35.78	40.37	29.5	*	+
	NT2RP4001828	9.76	9.76	34.37	27.78	34.21	26.3		
	NT2RP4001836	7.74	7.74	33.26	27.19	39.14	26.78		
15	NT2RP4001838	1.59	1.59	7.49	2.09	4.5	2.71		
	NT2RP4001841	8.75	8.75	80.37	61.67	50.27	56.46		
	NT2RP4001849	1.9	1.9	4.55	2.51	5.08	1.58		
	NT2RP4001861	7.27	7.27	34.6	36.09	41.9	34.39		
20	NT2RP4001877	6.59	6.59	8.44	12.87	9.04	14.1	*	- +
	NT2RP4001879	9.64	9.64	15.47	11.58	14.24	10.73		
	NT2RP4001889	5.09	5.09	10.66	6.68	11.25	8.91		
25	NT2RP4001893	3.97	3.97	7.34	3.19	6.11	2.72		
	NT2RP4001896	3.18	3.18	6.86	5.38	7.87	4.92		
	NT2RP4001898	7.83	7.83	26.41	22.98	20.13	20.15		
	NT2RP4001901	1.73	1.73	4.69	4.08	5.87	2.69		
30	NT2RP4001910	39.51	39.51	58.21	53.45	33.59	45.93		
	NT2RP4001925	4.1	4.1	10.21	6.69	6.32	6.12		
	NT2RP4001926	6.41	6.41	7.22	7.1	9.54	5.52		
35	NT2RP4001927	6.26	6.26	9.97	5.83	8.13	2.82		
	NT2RP4001931	8.64	8.64	14.16	15.49	17.54	11.89		
	NT2RP4001933	38.49	38.49	87.13	96.81	133.51	91.22		
	NT2RP4001938	2.93	2.93	4.53	4.27	3.31	3.55		
40	NT2RP4001942	13.44	13.44	27.12	31.34	24.8	18.71		
	NT2RP4001945	1.41	1.41	4	1.55	3.67	1.77		
	NT2RP4001946	1.97	1.97	5.67	3.04	3.96	1.66		
45	NT2RP4001947	4.42	4.42	8.93	5.92	9.81	5.16		
	NT2RP4001950	4.13	4.13	5	2.25	2.84	1.34	*	-
	NT2RP4001953	10.44	10.44	14.15	13.81	19.4	14.36		
	NT2RP4001966	2.44	2.44	2.41	2.51	4.26	1.52		
50	NT2RP4001970	2.26	2.26	5.32	3.91	3.4	2.88		
	NT2RP4001975	8.56	8.56	20.03	18.32	13.05	12.02		
	NT2RP4001988	6.72	6.72	18.78	22.92	24.78	29.44	*	+
	NT2RP4001996	5.27	5.27	12.83	10.65	16.35	12.42		
55	NT2RP4002014	3.4	3.4	8.14	43.19	37.87	33.17	**	+

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	NT2RP4001568	7.58	7.58	11.65	10.02	19.33	12.97		
	NT2RP4001569	15.49	15.49	41.42	32.56	40.11	27.18		
5	NT2RP4001571	5.75	5.75	13.28	9.06	7.43	8.13		
	NT2RP4001574	8.5	8.5	19.03	15.54	17.52	15.29		
	NT2RP4001575	2.46	2.46	5.86	5.71	5.53	4.24		
10	NT2RP4001578	17.21	17.21	45.64	52.76	53.63	40.92		
	NT2RP4001592	9.76	9.76	13.68	14.13	19.64	11.68		
	NT2RP4001593	9.4	9.4	18.11	22.12	27.3	18.04		
	NT2RP4001605	5.97	5.97	4.78	4.1	7.77	4.66		
15	NT2RP4001606	2.9	2.9	8.34	6.01	3.51	6.75		
	NT2RP4001607	2.04	2.04	5.24	4.57	3.26	3.78		
	NT2RP4001610	1.74	1.74	2.42	2.6	2.48	1.7		
20	NT2RP4001614	2.17	2.17	7.19	5.38	4.34	6.86		
	NT2RP4001623	2.38	2.38	5.26	2.43	2.65	2.02		
	NT2RP4001626	9.48	9.48	11.67	18.67	23.9	19.44	**	+
	NT2RP4001634	2.74	2.74	4.93	3.67	5.24	4.26		
25	NT2RP4001638	3.41	3.41	3.03	2.36	2.11	1.87	**	-
	NT2RP4001644	7.86	7.86	33.73	24.36	26.04	24.99		
	NT2RP4001646	11.61	11.61	15.02	7.42	10.74	11.21		
	NT2RP4001656	3.75	3.75	5.23	2.89	4.51	2.07		
30	NT2RP4001666	1.99	1.99	4.68	3.26	6.25	2.02		
	NT2RP4001670	11.74	11.74	15.51	12.45	7.09	8.31		
	NT2RP4001677	28.27	28.27	42.75	42.01	45.48	47.53		
35	NT2RP4001679	8.82	8.82	33.83	33.23	51.5	29.78		
	NT2RP4001695	7.71	7.71	12.76	15.66	20.35	12.87		
	NT2RP4001696	2.64	2.64	5.45	3.13	3.92	3.72		
	NT2RP4001699	3.58	3.58	8.03	3.18	4.12	4.42		
40	NT2RP4001717	2.79	2.79	7.03	3.29	5.84	4.15		
	NT2RP4001719	3.59	3.59	9.11	7.6	9.27	6.28		
	NT2RP4001725	2.27	2.27	4.79	2.28	5.07	1.43		
45	NT2RP4001726	7.07	7.07	11.18	5.85	6.91	4.98		
	NT2RP4001730	3.11	3.11	12.82	11.96	19.81	16.3		
	NT2RP4001739	2.83	2.83	5.83	5.79	6.55	4.98		
	NT2RP4001741	7.25	7.25	15.93	9.28	12.42	10.97		
50	NT2RP4001753	3.04	3.04	8.4	4.39	4.64	6.64		
	NT2RP4001760	4.32	4.32	6.6	7.79	7.73	12.96		
	NT2RP4001787	67.61	67.61	173.05	169.17	187.1	193.22		
55	NT2RP4001790	2	2	5.29	3.42	2.97	2.58		
	NT2RP4001795	9.31	9.31	12.31	14.38	19.76	12.34		

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	NT2RP4001373	8.86	8.86	16.4	10.99	8.59	11.48		
	NT2RP4001375	2.71	2.71	6.06	4.91	7.42	2.94		
5	NT2RP4001379	1.74	1.74	3.52	2.34	4.67	1.38		
	NT2RP4001381	5.6	5.6	12.51	12.8	12.66	8.27		
	NT2RP4001386	6.39	6.39	14.52	14.77	20.11	12.32		
	NT2RP4001389	7.28	7.28	8.66	5.43	9.25	4.09		
10	NT2RP4001396	5.76	5.76	6.42	2.83	4.61	2.12	*	-
	NT2RP4001407	2.92	2.92	2.98	3.07	2.04	1.76		
	NT2RP4001409	13.6	13.6	28.28	8.87	5.85	8.84		
15	NT2RP4001410	33.56	33.56	61.26	40.57	18.92	37.8		
	NT2RP4001414	16.59	16.59	37.89	14.29	21.3	16.8		
	NT2RP4001424	3.55	3.55	8.85	7.99	10.45	7.41		
	NT2RP4001433	3.85	3.85	6	8.39	9.79	3.38		
20	NT2RP4001438	9.95	9.95	27.94	46.22	53.63	30.76	*	+
	NT2RP4001442	4.33	4.33	4.97	3.3	3.41	2.64	*	-
	NT2RP4001447	4.42	4.42	4.69	5.08	5.51	3.41		
25	NT2RP4001466	3.74	3.74	5.45	7.38	3.23	5.57		
	NT2RP4001467	21.67	21.67	58.89	54.18	44.07	55.8		
	NT2RP4001472	8.05	8.05	12.19	11.7	11.76	11.97		
	NT2RP4001474	2.83	2.83	4.81	3.2	5.73	1.98		
30	NT2RP4001483	1.48	1.48	4.33	2.61	3.7	1.19		
	NT2RP4001488	21.03	21.03	32.41	39.07	46.9	33.42		
	NT2RP4001492	4.18	4.18	6.73	3.21	3.81	1.91		
35	NT2RP4001498	4.3	4.3	2.43	2.95	3.52	2.13		
	NT2RP4001502	28.2	28.2	57.38	27.65	34.81	41.83		
	NT2RP4001503	3.83	3.83	6.74	5.51	4.1	3.37		
	NT2RP4001507	2.39	2.39	3.69	5.03	5.95	5.18	**	+
40	NT2RP4001510	1.74	1.74	4.63	5.64	5.05	3.02		
	NT2RP4001516	3.54	3.54	4.16	2.52	1.9	1.27	*	-
	NT2RP4001520	25.33	25.33	70.64	73.93	107.21	85.05		
45	NT2RP4001523	5.57	5.57	9.99	8.4	7.79	6.19		
	NT2RP4001524	6.1	6.1	10.17	8.92	11.45	5.98		
	NT2RP4001529	29.8	29.8	74.89	59.39	60.69	55.29		
	NT2RP4001531	2.88	2.88	10.96	8.63	11.05	7.81		
50	NT2RP4001546	143.48	143.48	388.31	316.63	215.6	309.45		
	NT2RP4001547	9.05	9.05	26.31	28.76	23.06	18.81		
	NT2RP4001551	2.27	2.27	3.93	3.87	4.08	2.02		
	NT2RP4001555	2.95	2.95	5.8	4.14	3.58	1.74		
55	NT2RP4001567	6.54	6.54	6.88	5.82	7.12	5.68		

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	NT2RP4001159	7.08	7.08	11.61	7.69	5.58	10.46		
	NT2RP4001162	3.77	3.77	6.14	4.07	6.06	3.41		
5	NT2RP4001170	1.15	1.15	4.53	1.28	3.2	2.15		
	NT2RP4001174	4.16	4.16	12.27	7.91	11.95	5.02		
	NT2RP4001175	9.65	9.65	19.14	15.72	21.29	13.28		
10	NT2RP4001176	99.19	99.19	161.51	174.03	241.92	194	*	+
	NT2RP4001184	4.83	4.83	27.1	25.76	32.51	22.85		
	NT2RP4001198	21.66	21.66	48.22	29.54	29.17	38.54		
	NT2RP4001199	2.52	2.52	6.45	2.59	4.74	3.71		
15	NT2RP4001206	8.25	8.25	33.2	25.92	32.07	25.48		
	NT2RP4001207	2.38	2.38	5.15	2.21	3.11	4.01		
	NT2RP4001210	2.73	2.73	5.2	3.62	4.26	2.64		
	NT2RP4001213	3.42	3.42	5.11	3.99	4.23	3.63		
20	NT2RP4001214	3.34	3.34	4.3	3.51	3.76	2.16		
	NT2RP4001219	7.4	7.4	12.05	14.35	19.28	13.39	*	+
	NT2RP4001228	5.26	5.26	9.63	12.15	15.74	20.07	*	+
25	NT2RP4001235	2.42	2.42	7.45	3.46	6.02	4.48		
	NT2RP4001256	2.11	2.11	4.24	1.66	3.41	2.66		
	NT2RP4001257	2.48	2.48	7.27	4.05	4.35	4.05		
	NT2RP4001260	3.16	3.16	5.79	2.52	3.86	2.92		
30	NT2RP4001261	3.84	3.84	6.63	8.42	6.47	5		
	NT2RP4001274	22.92	22.92	38.08	25.02	31.56	21.25		
	NT2RP4001276	5.24	5.24	10.03	11.38	15.97	11.63	*	+
35	NT2RP4001283	20.72	20.72	122.55	87.44	93.43	86.47		
	NT2RP4001299	9.62	9.62	15.14	14.95	10.52	15.18		
	NT2RP4001313	1.45	1.45	3.26	1.72	2.96	0.97		
	NT2RP4001315	6.06	6.06	11.14	7.45	9.92	7.74		
40	NT2RP4001320	14.6	14.6	42.74	32.02	38.13	29.24		
	NT2RP4001325	32.53	32.53	146.14	142.88	178.36	128.89		
	NT2RP4001336	6.69	6.69	40.75	38.55	46.66	32.11		
45	NT2RP4001339	4.12	4.12	5.6	3.35	5.56	2.76		
	NT2RP4001343	10.46	10.46	83.37	54.71	61.01	60.06		
	NT2RP4001344	6.7	6.7	60.08	49.79	55.21	42.62		
	NT2RP4001345	1.65	1.65	6.68	5.64	5.7	3.87		
50	NT2RP4001351	4.1	4.1	15.97	10.01	20.05	11.42		
	NT2RP4001353	2.8	2.8	5.91	1.63	2.94	1.86		
	NT2RP4001355	2.57	2.57	8.67	1.83	3.12	2.08		
55	NT2RP4001367	10.64	10.64	17.66	11.92	17.06	13.16		
	NT2RP4001372	2.26	2.26	3.82	2.1	2.06	2.07		

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	NT2RP4000996	3.54	3.54	7.49	6.77	6.52	7.38		
	NT2RP4000997	21.59	21.59	36.81	28.52	15.18	34.38		
5	NT2RP4001001	5.53	5.53	9.17	16.66	18.38	15.09	**	+
	NT2RP4001004	1.71	1.71	4.88	2.84	3.09	1.37		
	NT2RP4001006	3.46	3.46	8.12	6.85	6.52	6.13		
10	NT2RP4001009	9.3	9.3	10.45	15.44	20.46	8.25		
	NT2RP4001010	7.33	7.33	9.13	7.38	9.75	6.68		
	NT2RP4001013	23.29	23.29	50.16	30.87	28.1	30.91		
	NT2RP4001029	2.49	2.49	5.95	4.05	2.84	3.63		
15	NT2RP4001036	7.55	7.55	13.55	9.11	11.51	13.16		
	NT2RP4001041	6.57	6.57	14.4	9.89	12.3	6.35		
	NT2RP4001042	4.34	4.34	8.11	9.44	12.5	8.79		
20	NT2RP4001046	6.98	6.98	9.95	13.24	16.28	15.36	**	+
	NT2RP4001050	5.28	5.28	4.81	3.79	4.64	3.35	*	-
	NT2RP4001051	6.48	6.48	8.44	5.43	6.82	5.26		
	NT2RP4001057	0.76	0.76	2.19	2.34	2.43	1.87		
25	NT2RP4001063	1.48	1.48	4.39	3.34	3.53	1.8		
	NT2RP4001064	3.51	3.51	9.18	12.02	9.13	11.57		
	NT2RP4001067	4.42	4.42	9.77	10.96	9.63	6.6		
	NT2RP4001078	2.12	2.12	3.43	2.67	2.53	1.82		
30	NT2RP4001079	5.3	5.3	9.35	8.51	8.02	8.98		
	NT2RP4001080	4.1	4.1	5.27	3.52	4.52	2.3		
	NT2RP4001086	5.08	5.08	4.19	3.93	6.64	2.85		
35	NT2RP4001095	2.49	2.49	7.25	7.96	6.49	6.85		
	NT2RP4001098	0.92	0.92	3.38	3.87	2.95	3.41		
	NT2RP4001100	6.47	6.47	24.34	20.89	20.64	16.99		
	NT2RP4001105	3.13	3.13	7.23	6.51	5.58	4.61		
40	NT2RP4001110	1.75	1.75	3.5	7.07	8.35	5.29	*	+
	NT2RP4001115	9.95	9.95	17.68	20.6	18.48	15.31		
	NT2RP4001117	19.81	19.81	30.49	35.35	42.53	27.5		
45	NT2RP4001122	6.06	6.06	6.09	5.17	6.25	3.27		
	NT2RP4001123	3.62	3.62	7.76	7.95	5.96	6.27		
	NT2RP4001126	4.36	4.36	11.28	10.87	9.09	8.04		
	NT2RP4001127	3.25	3.25	4.59	3.39	3.08	2.17		
50	NT2RP4001138	2.46	2.46	5.8	3.41	2.56	1.62		
	NT2RP4001143	2.73	2.73	5.98	6.44	6.54	5.66		
	NT2RP4001148	3.72	3.72	6.76	3.77	3.03	2.05		
55	NT2RP4001149	5.07	5.07	7.28	6.76	9.03	6.37		
	NT2RP4001150	3.8	3.8	3.17	3.15	3.7	2.88		

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	NT2RP4000839	12.23	12.23	97.13	79.71	85.74	86.06		
	NT2RP4000846	3.8	3.8	10.13	4.65	3.46	6.65		
5	NT2RP4000848	4.63	4.63	10.74	8.65	8.58	6.07		
	NT2RP4000855	2.91	2.91	4.7	4	3.85	3.43		
	NT2RP4000863	3.08	3.08	4.33	3.11	5.3	3.78		
10	NT2RP4000865	6.43	6.43	25.36	20.09	39.64	21.24		
	NT2RP4000873	9.64	9.64	88.25	63.22	69.65	71.33		
	NT2RP4000874	1.76	1.76	3.98	2.37	3.67	2.03		
	NT2RP4000875	3.31	3.31	9.24	6.88	6.52	7.19		
15	NT2RP4000878	24.17	24.17	42.53	28.01	16.35	29.04		
	NT2RP4000879	2.56	2.56	5.1	2.95	5.29	2.62		
	NT2RP4000880	5.17	5.17	21.59	20.97	27.22	16.8		
20	NT2RP4000891	81.07	81.07	192.57	252.29	351.53	221.08	*	+
	NT2RP4000894	5.16	5.16	9.81	8.53	4.8	6.97		
	NT2RP4000898	0.86	0.86	2.74	1.88	2.14	1.64		
	NT2RP4000899	9.63	9.63	29.48	24.01	20.85	23.95		
25	NT2RP4000907	2.14	2.14	3.58	1.74	4.04	0.81		
	NT2RP4000908	4.62	4.62	9.67	7.51	5.9	5.87		
	NT2RP4000910	14.4	14.4	104.68	124.04	197.74	160.9	*	+
	NT2RP4000918	2.85	2.85	4.76	4.73	4.26	5.35		
30	NT2RP4000925	3.9	3.9	5.53	2.81	3.15	1.86		
	NT2RP4000927	1.99	1.99	2.5	0.46	1.08	0.6	**	-
	NT2RP4000928	3.11	3.11	6.8	4.48	4.22	5.18		
35	NT2RP4000929	1.44	1.44	3.68	1.94	2.86	0.84		
	NT2RP4000946	0.92	0.92	3.41	1.78	3.22	1.47		
	NT2RP4000947	1.71	1.71	3.51	1.94	3.31	1.79		
	NT2RP4000949	4.94	4.94	7.84	3.88	5.48	2.38		
40	NT2RP4000955	4.17	4.17	6.34	2.07	2.86	0.54	*	-
	NT2RP4000959	20.55	20.55	28.14	36.21	42.82	36.14	*	+
	NT2RP4000962	3.4	3.4	4.24	8.33	10.09	4.53		
45	NT2RP4000973	8.6	8.6	16.04	10.31	8.92	9.03		
	NT2RP4000975	2.18	2.18	5.84	3.29	3.05	2.62		
	NT2RP4000979	3.83	3.83	8.67	5.55	8.13	7.1		
	NT2RP4000984	1.61	1.61	4.31	3.15	3.93	1.85		
50	NT2RP4000986	7.32	7.32	13.27	12.66	12.35	2.52		
	NT2RP4000988	5.74	5.74	8.37	4	5.2	2.63		
	NT2RP4000989	6.24	6.24	6.55	4.05	3.48	2.89	**	-
	NT2RP4000990	3.16	3.16	4	1.92	1.69	2.16	**	-
55	NT2RP4000994	4.04	4.04	7.67	16.48	10.13	15.95	*	+

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	NT2RP4000568	1.72	1.72	3.83	5.6	6.08	4.46	*	+
	NT2RP4000583	11.61	11.61	20.2	23.65	14.97	20.56		
5	NT2RP4000585	3.04	3.04	4.14	3.12	2.55	3.24		
	NT2RP4000588	8.65	8.65	12.77	14.58	16.96	13.55	*	+
	NT2RP4000590	24.89	24.89	41.97	41.86	50.81	32.65		
10	NT2RP4000599	4.29	4.29	3.24	2.44	2.23	3.59		
	NT2RP4000603	14.08	14.08	33.32	31.06	21.01	29.12		
	NT2RP4000607	2.41	2.41	10.04	4.45	7.87	15.35		
	NT2RP4000614	6.14	6.14	15.21	15.57	12.53	15.19		
15	NT2RP4000634	6.61	6.61	11	7.78	9.84	10.31		
	NT2RP4000638	3.59	3.59	7.77	8.45	5.8	4.73		
	NT2RP4000648	3.13	3.13	4.26	2.54	2.69	2.19		
20	NT2RP4000657	9.94	9.94	15.38	15.95	18.93	14.49		
	NT2RP4000691	5.76	5.76	5.82	4.92	7.47	7.73		
	NT2RP4000697	3.74	3.74	8.5	5.55	6.56	6.12		
	NT2RP4000704	8.91	8.91	47.2	44.17	54.81	38.14		
25	NT2RP4000710	40.22	40.22	98.85	90.4	59.28	83.71		
	NT2RP4000713	4.35	4.35	19.92	16.67	20.85	15.52		
	NT2RP4000724	6.29	6.29	12.5	8.19	9.81	7.83		
30	NT2RP4000725	3.61	3.61	4	1.88	1.74	2.33	**	-
	NT2RP4000728	10.13	10.13	41.12	43.53	66.46	39.83		
	NT2RP4000737	4.07	4.07	2.15	3.63	3.09	3.28		
	NT2RP4000739	5.07	5.07	7.71	4.61	3.63	5.84		
35	NT2RP4000749	2.4	2.4	5.29	2.59	3.97	1.68		
	NT2RP4000769	4.93	4.93	10.12	4.67	6.27	6.2		
	NT2RP4000774	3.34	3.34	8.87	5.12	6.63	4.27		
	NT2RP4000781	2.15	2.15	5.12	2.06	2.26	1.55		
40	NT2RP4000783	6.81	6.81	15.16	13.48	15.44	12.67		
	NT2RP4000787	1.45	1.45	2.27	0.31	0.51	0.54	*	-
	NT2RP4000788	3.58	3.58	23.26	16	18.3	18.73		
45	NT2RP4000792	3.68	3.68	5.64	5.5	5.8	9.45		
	NT2RP4000809	43.7	43.7	56.09	46.75	50.47	81.62		
	NT2RP4000817	3.65	3.65	7.83	7.92	7.25	5.82		
	NT2RP4000821	31.34	31.34	38.66	28.32	33.11	25.22		
50	NT2RP4000822	2.46	2.46	5.91	4.29	6.19	2.6		
	NT2RP4000823	697.74	697.74	1127.48	923.16	1026.8	947.85		
	NT2RP4000831	9.98	9.98	61.97	44.37	68.47	50.69		
55	NT2RP4000833	3.19	3.19	11.26	6.73	7.19	11.91		
	NT2RP4000837	1.41	1.41	4.03	1.56	3.65	2.29		

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	NT2RP4000447	10.3	10.3	64.21	64.91	71.82	68.48		
	NT2RP4000448	5.59	5.59	6.67	4.32	3.56	6.27		
5	NT2RP4000449	5.67	5.67	-20.48	14.2	19.45	14.67		
	NT2RP4000453	3.53	3.53	6.32	9.02	10.32	10.12	**	+
	NT2RP4000455	2.64	2.64	3.98	1.81	1.75	2.4		
	NT2RP4000456	12.57	12.57	21.7	14.7	9.86	13.58		
10	NT2RP4000457	1.54	1.54	7.12	3.9	7.55	2.98		
	NT2RP4000461	5.7	5.7	9.84	7.65	6.41	4.6		
	NT2RP4000462	11.76	11.76	15.32	11.86	11.53	17.37		
15	NT2RP4000463	10.2	10.2	52.59	50.66	69.3	48.36		
	NT2RP4000471	6.36	6.36	10.74	5.74	6.23	4.98		
	NT2RP4000472	3.97	3.97	4.41	1.27	1.27	1.66	**	-
	NT2RP4000476	27.14	27.14	52.56	74.95	94.93	65.35	*	+
20	NT2RP4000480	11.56	11.56	29.27	19.08	9.95	26.66		
	NT2RP4000481	2.29	2.29	4.76	3.73	4.16	4.33		
	NT2RP4000483	1.38	1.38	7.59	7.58	7.85	6.26		
25	NT2RP4000487	1.54	1.54	4.9	2.26	3.17	0.91		
	NT2RP4000496	2.16	2.16	4.98	2.95	4.07	3.65		
	NT2RP4000497	17.15	17.15	22.33	34.9	44.46	29.9	*	+
	NT2RP4000498	10.46	10.46	21.39	20.15	30.33	24.49		
30	NT2RP4000500	2.43	2.43	3.21	2.03	1.49	1.39	*	-
	NT2RP4000507	5.63	5.63	9.02	12.49	10.62	17.51	*	+
	NT2RP4000515	12.85	12.85	88.89	90.3	101.29	96.12		
35	NT2RP4000516	8.77	8.77	26.09	19.63	21.76	21.82		
	NT2RP4000517	3.22	3.22	5.73	4.69	5.89	4.79		
	NT2RP4000518	3.42	3.42	7.4	4.47	6.05	3.82		
	NT2RP4000519	2.9	2.9	5.76	2.18	2.22	1.77		
40	NT2RP4000524	4.2	4.2	3.44	2.05	1.72	1.3	**	-
	NT2RP4000528	3.67	3.67	3.06	3.01	3.27	8.01		
	NT2RP4000537	35.4	35.4	62.6	36.23	30.93	44.52		
45	NT2RP4000541	2.04	2.04	2.45	3.34	4.33	3.09	*	+
	NT2RP4000543	2.93	2.93	8.94	7.96	9.72	8.75		
	NT2RP4000545	4.03	4.03	6.38	4.99	6.43	4.78		
	NT2RP4000546	3.34	3.34	5.93	5.53	4.9	6.03		
50	NT2RP4000549	23.81	23.81	56.48	41.6	51.57	38.82		
	NT2RP4000556	7.36	7.36	13.04	14.69	15.54	12.07		
	NT2RP4000557	6.1	6.1	4.53	1.82	3.97	6.27		
55	NT2RP4000558	30.12	30.12	94.28	68.16	57.01	73.2		
	NT2RP4000560	14.8	14.8	52.31	49.75	58.69	56.12		

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	NT2RP4000250	16.86	16.86	31.23	24.92	23.85	26.39		
	NT2RP4000256	4.38	4.38	8.45	4.99	6.44	4.67		
5	NT2RP4000257	32.45	32.45	44.22	48.67	57.52	39.19		
	NT2RP4000259	7.07	7.07	6.96	8.48	13.07	9.08		
	NT2RP4000261	4.18	4.18	8.13	4.07	4.55	6.48		
	NT2RP4000262	8.36	8.36	11.02	7.11	10.05	7.64		
10	NT2RP4000263	2.6	2.6	5.14	3.47	3.67	3.78		
	NT2RP4000280	5.84	5.84	11.62	9.05	11.58	14.23		
	NT2RP4000286	4.73	4.73	10.16	4.38	6.77	11.94		
15	NT2RP4000290	5.77	5.77	5.42	3.18	3.19	5.5		
	NT2RP4000291	42.53	42.53	73.59	62.12	70.23	70.61		
	NT2RP4000301	3.31	3.31	20.59	22.93	34.62	26.63	*	+
	NT2RP4000312	7.76	7.76	45.27	39.01	43.59	45.92		
20	NT2RP4000321	3.73	3.73	8.16	7.88	7.14	9.75		
	NT2RP4000323	1.44	1.44	4.26	2.27	2.55	2.12		
	NT2RP4000324	7.77	7.77	16.76	8.33	10.2	11.08		
25	NT2RP4000334	20.97	20.97	77.78	63.81	71.76	68.24		
	NT2RP4000343	2.48	2.48	5.54	1.57	3.16	2.38		
	NT2RP4000348	3.4	3.4	10.81	8.38	6.75	13.2		
	NT2RP4000349	1.78	1.78	0.83	0.37	0.72	2.48		
30	NT2RP4000355	3.98	3.98	14.84	7.18	9.13	9.04		
	NT2RP4000356	8.3	8.3	22.64	9.22	11.44	21.13		
	NT2RP4000360	3.04	3.04	6.98	5.06	5.43	4.73		
35	NT2RP4000367	2.18	2.18	3.72	2.04	2.58	2.83		
	NT2RP4000370	4.21	4.21	7.51	3.62	7.12	5.99		
	NT2RP4000373	3.33	3.33	5.18	3.19	5.23	2.95		
	NT2RP4000376	4.2	4.2	5.5	4.85	5.53	6.5		
40	NT2RP4000381	1.97	1.97	6.46	4.31	7.01	5.59		
	NT2RP4000388	85.82	85.82	204.63	128.93	93.38	116.03		
	NT2RP4000390	12.16	12.16	94.1	78.97	76.6	82.02		
45	NT2RP4000393	8.66	8.66	9.77	9.09	5.99	10.79		
	NT2RP4000398	5.51	5.51	26.52	22.08	28.3	26.61		
	NT2RP4000406	6.68	6.68	15.61	12.95	15.02	10.08		
	NT2RP4000407	6.17	6.17	11.52	9.41	14.74	9.13		
50	NT2RP4000413	1.79	1.79	3.63	1.15	2.35	2.12		
	NT2RP4000415	7.59	7.59	26.11	18.6	21.68	21.68		
	NT2RP4000417	7.76	7.76	26.64	14.47	12.79	19.19		
	NT2RP4000423	3.52	3.52	7.56	7	6.98	6.46		
55	NT2RP4000424	2.51	2.51	7.2	3.07	5.03	3.27		

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	NT2RP4000065	7.54	7.54	9.24	11.85	15.01	11.47	*	+
	NT2RP4000070	6.63	6.63	5.29	3.79	4.54	3.84	*	-
5	NT2RP4000074	6.55	6.55	43.63	10.29	15.39	9.52		
	NT2RP4000078	3.41	3.41	9.55	12.2	11.66	12.56	*	+
	NT2RP4000080	3.52	3.52	7.01	5.06	6.84	6.81		
10	NT2RP4000099	128.86	128.86	236.22	149.44	161.1	211.56		
	NT2RP4000102	3.55	3.55	6.48	5.27	4.97	5.54		
	NT2RP4000103	2	2	5.47	2.57	2.85	3.96		
	NT2RP4000108	4.66	4.66	7.91	9.33	10.73	10.61	*	+
15	NT2RP4000109	18.89	18.89	22.84	19.87	24.15	16.14		
	NT2RP4000111	4.56	4.56	6.38	4.87	5.27	4.85		
	NT2RP4000112	5.62	5.62	10.14	12.45	8.7	11.81		
20	NT2RP4000115	2.94	2.94	3.62	4.95	7.82	10.93		
	NT2RP4000129	2.18	2.18	4.58	3.02	5.04	5.03		
	NT2RP4000137	3.36	3.36	10.05	5.34	8.72	11.4		
	NT2RP4000138	7.21	7.21	10.91	17.75	20.19	19.17	**	+
25	NT2RP4000141	3.25	3.25	6.1	5.22	4.9	3.57		
	NT2RP4000147	6.21	6.21	4.49	4.27	5.2	3.47		
	NT2RP4000150	5.96	5.96	6.93	7.33	10.41	7.06		
30	NT2RP4000151	2.82	2.82	7.82	5.69	6.24	5.05		
	NT2RP4000157	73.27	73.27	222.87	169.53	97.5	173.8		
	NT2RP4000159	2.02	2.02	5.03	3.38	5.03	2.92		
	NT2RP4000163	5.21	5.21	8.74	10.89	9.18	6.63		
35	NT2RP4000167	3.26	3.26	4.35	4.32	3.22	3.69		
	NT2RP4000171	5.72	5.72	7	5.72	7.45	5.03		
	NT2RP4000175	62.48	62.48	94.56	144.06	214.73	147.88	*	+
40	NT2RP4000180	17.17	17.17	106.15	88.61	123.6	81.8		
	NT2RP4000185	7.64	7.64	31.76	31.82	23.23	34.01		
	NT2RP4000192	1.04	1.04	4.78	4.19	3.6	3.05		
	NT2RP4000194	3.13	3.13	7.53	7.58	4.98	6.36		
45	NT2RP4000196	6.81	6.81	43.94	35.57	46.56	41.91		
	NT2RP4000210	5.63	5.63	9.71	9.96	8.35	9.27		
	NT2RP4000212	5.59	5.59	8.88	8.57	8.41	9.08		
	NT2RP4000214	5.53	5.53	10.21	5.72	6.68	11.29		
50	NT2RP4000216	8.89	8.89	7.36	7	12.63	10.08		
	NT2RP4000218	3.45	3.45	9.78	7.25	6.07	5.77		
	NT2RP4000223	21.18	21.18	177.28	121.8	125.75	125.3		
55	NT2RP4000243	16.52	16.52	54.51	42.94	41.61	51.74		
	NT2RP4000246	17.75	17.75	37.97	26.43	11.42	27.5		

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	NT2RP3004524	10.04	10.04	29.21	26.88	26.43	29.62		
	NT2RP3004527	3.03	3.03	3.22	1.29	2.23	2.08	*	-
5	NT2RP3004534	3.08	3.08	10.44	5.58	12	7.38		
	NT2RP3004539	4.45	4.45	12.17	9.8	8.97	12		
	NT2RP3004541	2.65	2.65	11.06	5.48	9.44	8.25		
10	NT2RP3004544	3.54	3.54	8.89	6.62	5.24	9.48		
	NT2RP3004551	3.46	3.46	6.75	6.6	6.6	6.98		
	NT2RP3004552	2.76	2.76	4.33	2.84	3.22	4.98		
	NT2RP3004557	5.68	5.68	8.73	9.74	15.44	13.16	*	+
15	NT2RP3004561	1.96	1.96	3.77	2.55	4.46	4.1		
	NT2RP3004566	3.09	3.09	11.55	7.2	10.05	8.79		
	NT2RP3004569	2.21	2.21	7.09	4.63	5.36	6.91		
20	NT2RP3004572	4.37	4.37	6.83	7.08	5.55	7.07		
	NT2RP3004578	2.35	2.35	5.38	4.15	4.27	3.24		
	NT2RP3004584	4.76	4.76	28.36	34.99	37.13	30.29		
	NT2RP3004588	2.38	2.38	4.89	1.6	3.7	3.28		
25	NT2RP3004594	2.25	2.25	5.9	5.67	6.49	8.94		
	NT2RP3004603	34.16	34.16	99.64	80.2	102.6	97.27		
	NT2RP3004612	4.71	4.71	12.17	5.3	3.36	5.34		
30	NT2RP3004617	1.09	1.09	2.32	2.49	3.3	2.39		
	NT2RP3004618	4.61	4.61	5.9	2.49	5.21	5.9		
	NT2RP3004625	3.97	3.97	8.17	4.55	6.92	7.1		
	NT2RP3004635	4.76	4.76	7.83	1.52	2.86	3.47		
35	NT2RP3004640	10.61	10.61	62.15	59.33	67.97	48.32		
	NT2RP3004642	8.04	8.04	29.31	22.82	26.12	25.12		
	NT2RP3004647	3.5	3.5	5.65	5.89	7.35	6.88	*	+
	NT2RP3004652	1.76	1.76	10.37	4.2	3.71	4.34		
40	NT2RP3004669	2.01	2.01	5.36	4.01	5.33	3.46		
	NT2RP3004670	5.04	5.04	10.58	12.4	9.19	14.23		
	NT2RP4000008	45.17	45.17	71.24	49.77	32.43	48.77		
45	NT2RP4000018	11.64	11.64	14.61	11.69	14.8	14.87		
	NT2RP4000023	6.96	6.96	8.91	4.86	7.38	5.98		
	NT2RP4000025	16.2	16.2	22.16	26.22	29.89	24.7	*	+
	NT2RP4000035	6.3	6.3	12.01	11.28	15.33	11.01		
50	NT2RP4000041	14.46	14.46	34.8	22.01	17.41	23.68		
	NT2RP4000049	2.64	2.64	6.34	6.59	6.88	5.3		
	NT2RP4000050	2.24	2.24	6.87	3.54	4.48	4.05		
	NT2RP4000051	4.66	4.66	10.5	10.58	10.02	8.96		
55	NT2RP4000063	20.51	20.51	33.5	26.77	31	18.33		

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	NT2RP3004332	11.69	11.69	100.11	78.54	102.41	76.72		
	NT2RP3004334	1.49	1.49	6.97	8.56	6.06	9.06		
5	NT2RP3004336	2.11	2.11	-6.24	6.02	4.44	5.63		
	NT2RP3004338	3.09	3.09	8.41	10.22	9.52	16.47		
	NT2RP3004341	1.81	1.81	4.56	6.13	5.17	9.13		
10	NT2RP3004345	4.1	4.1	8.68	9.3	9.63	8.48		
	NT2RP3004348	5.06	5.06	11.25	13.04	10.79	12.54		
	NT2RP3004349	5	5	7.5	4.89	7.75	5.76		
	NT2RP3004355	5.57	5.57	7.09	7.55	7.07	7.18		
15	NT2RP3004356	5.76	5.76	21.51	11.29	15.14	15.56		
	NT2RP3004360	3.4	3.4	5.26	6.01	5.32	7.85		
	NT2RP3004361	2.6	2.6	6.26	7.67	7.3	8.87	*	+
	NT2RP3004374	3.06	3.06	10.09	8.8	6.6	5.75		
20	NT2RP3004378	10.48	10.48	18.57	28.26	24.09	34.81	*	+
	NT2RP3004399	3.88	3.88	5.77	3.53	3.17	9.06		
	NT2RP3004405	4.07	4.07	6.77	3.03	5.52	3.93		
25	NT2RP3004406	5.36	5.36	6.23	5.19	6.03	6.12		
	NT2RP3004411	5.93	5.93	13.28	8.08	6.39	9.51		
	NT2RP3004424	1.53	1.53	2.43	3.27	1.81	2.83		
	NT2RP3004428	3.03	3.03	5.36	5.07	3.82	4.09		
30	NT2RP3004432	3.3	3.3	3.52	3.61	3.11	4.38		
	NT2RP3004434	3.42	3.42	8.41	7.28	9.09	7.99		
	NT2RP3004446	3.29	3.29	4.6	3.29	4.1	2.63		
35	NT2RP3004451	3.2	3.2	6.01	3.89	3.38	2.48		
	NT2RP3004454	2.96	2.96	4.16	2.69	3.5	2.5		
	NT2RP3004466	3.5	3.5	7.89	5.25	3.85	5.61		
	NT2RP3004470	7.42	7.42	24.53	18.4	16.35	24.72		
40	NT2RP3004472	2.49	2.49	4.4	3.97	3.84	3.88		
	NT2RP3004475	1.71	1.71	5.52	2.72	5.93	3.9		
	NT2RP3004480	14.12	14.12	17.04	18.94	10.5	18.82		
45	NT2RP3004481	5.42	5.42	11.37	5.04	7.37	12.39		
	NT2RP3004490	2.66	2.66	8.45	3.92	7.03	8.25		
	NT2RP3004496	4.8	4.8	14.38	7.22	9.3	11.08		
	NT2RP3004498	6.39	6.39	21.39	16.86	15.11	18.92		
50	NT2RP3004503	-2.78	2.78	9.34	4.85	6.23	5.88		
	NT2RP3004504	3.91	3.91	11.09	6.05	10.67	11.52		
	NT2RP3004505	17.38	17.38	28.56	37.35	26.72	37.82		
	NT2RP3004507	1.57	1.57	6.6	2.52	5.52	1.68		
55	NT2RP3004519	4.9	4.9	7.73	4.93	8.36	8.75		

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	NT2RP3004093	2.38	2.38	7.49	4.07	3.51	4.47		
	NT2RP3004095	5.02	5.02	13.11	11.57	10.17	18.55		
5	NT2RP3004102	3.32	3.32	5.59	5.25	4.27	3.21		
	NT2RP3004110	12.74	12.74	18.66	22.12	14.31	19.97		
	NT2RP3004119	3.3	3.3	7.71	3.91	4.08	3.73		
	NT2RP3004125	5.55	5.55	12.05	8.13	10.88	8.38		
10	NT2RP3004129	4.62	4.62	7.38	3.36	2.95	6.08		
	NT2RP3004130	11.81	11.81	28.12	21.92	31.13	21.05		
	NT2RP3004133	4.51	4.51	12.95	14.62	8.94	16.81		
15	NT2RP3004145	1.43	1.43	4.17	2.62	4.87	3.59		
	NT2RP3004148	2.67	2.67	7.07	5.26	6.24	4.5		
	NT2RP3004155	2.37	2.37	4.82	4.7	4.57	6.59		
	NT2RP3004165	17.94	17.94	29.96	29.58	31.82	39.18		
20	NT2RP3004179	7.34	7.34	6.72	2.71	5.41	3.3	*	-
	NT2RP3004185	5.2	5.2	5.53	2.76	2.76	1.95	**	-
	NT2RP3004188	4.77	4.77	10.82	7.74	11.35	7		
25	NT2RP3004189	4.23	4.23	5.91	4.97	4.82	6.28		
	NT2RP3004190	2.6	2.6	5.57	5.84	4.36	5.26		
	NT2RP3004191	14.09	14.09	23.4	31.41	29.45	30.09	*	+
	NT2RP3004202	2.04	2.04	4.56	4.16	4.42	2.3		
30	NT2RP3004205	8.75	8.75	21.54	21.27	25.35	20.28		
	NT2RP3004206	4.5	4.5	9.74	5.14	6.37	9		
	NT2RP3004207	5.19	5.19	4.99	3.09	3.25	1.77	**	-
35	NT2RP3004209	4.74	4.74	7.74	8.2	11.23	9.08	*	+
	NT2RP3004215	1.86	1.86	6.7	3.96	2.41	4.55		
	NT2RP3004219	5.15	5.15	11.25	10.04	8.81	13.65		
	NT2RP3004242	4.65	4.65	10.36	9.8	10.19	14.56		
40	NT2RP3004246	4.5	4.5	9.39	9.18	10.95	3.8		
	NT2RP3004253	1.89	1.89	4.85	3.64	4.99	2.8		
	NT2RP3004258	5.45	5.45	10.89	12.77	11.07	11.39		
45	NT2RP3004262	4.26	4.26	5.71	2.63	2.99	2.01	*	-
	NT2RP3004275	5.59	5.59	3.43	1.4	2.97	2.34	*	-
	NT2RP3004282	5.45	5.45	68.08	51.29	52.72	53.57		
	NT2RP3004289	1.79	1.79	2.95	1.9	2.18	3.99		
50	NT2RP3004294	2.74	2.74	6.02	6.95	6.93	7.24	*	+
	NT2RP3004298	8.76	8.76	48.63	46.33	60.89	50.83		
	NT2RP3004309	3.3	3.3	6.46	5.2	5.22	5.31		
55	NT2RP3004321	3.71	3.71	6.11	3.29	3.74	3.34		
	NT2RP3004322	5.61	5.61	6.86	6.06	6.43	6.56		

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	NT2RP3003924	2.25	2.25	9.57	6.49	5.34	6.91		
	NT2RP3003932	1.41	1.41	3.85	5.17	5.26	3.85		
5	NT2RP3003939	3.48	3.48	11.88	9.86	14.05	11.09		
	NT2RP3003940	11.34	11.34	27.33	23.54	20.59	23.06		
	NT2RP3003943	2.6	2.6	2.83	2.85	2.78	3.88		
10	NT2RP3003959	3.52	3.52	6.96	6.54	5.93	5.49		
	NT2RP3003963	4.83	4.83	7.59	4.01	4.61	2.52		
	NT2RP3003965	11.14	11.14	13.85	18.75	20.67	17.22	**	+
	NT2RP3003972	26.1	26.1	40.32	22.13	17.98	38.48		
15	NT2RP3003973	2.85	2.85	4.33	1.96	3.6	3.96		
	NT2RP3003979	5.89	5.89	12.53	6.92	8.49	8.84		
	NT2RP3003980	3.52	3.52	9.41	9.34	8.89	7.92		
20	NT2RP3003982	4.2	4.2	4.63	2.44	1.6	4.61		
	NT2RP3003989	6.24	6.24	4.69	9.61	5.62	16.05		-
	NT2RP3003992	2.13	2.13	4.89	2.47	5.12	4.8		
	NT2RP3004000	2.81	2.81	6	1.72	3.22	2.62		
25	NT2RP3004001	11.38	11.38	19.94	11.62	11.37	21.11		
	NT2RP3004005	2.89	2.89	7.79	4.7	4.48	6.84		
	NT2RP3004013	2.23	2.23	7.2	2.66	4.87	3.57		
30	NT2RP3004016	1.5	1.5	7.1	2.22	3.14	2.88		
	NT2RP3004025	4.02	4.02	7.69	7.48	12.19	9.01		
	NT2RP3004030	7.05	7.05	12.64	13.97	15.8	17.66	*	+
	NT2RP3004041	5.65	5.65	11.38	10.48	9.57	19.81		
35	NT2RP3004042	15.22	15.22	102.33	97.27	103.6	99.67		
	NT2RP3004044	2.13	2.13	6.51	5.14	7.21	4.22		
	NT2RP3004051	2.6	2.6	5.79	2.23	5.51	4.69		
	NT2RP3004052	7.1	7.1	11.22	5.63	4.98	9.78		
40	NT2RP3004053	15.87	15.87	35.04	23.12	40.67	40.17		
	NT2RP3004055	2.38	2.38	5.33	2.98	3.3	4.47		
	NT2RP3004059	4.05	4.05	8.8	8.15	7.03	11		
45	NT2RP3004063	5.13	5.13	11.23	8.78	11.27	12.33		
	NT2RP3004067	4.24	4.24	8.4	6.62	6.42	4.47		
	NT2RP3004070	3.58	3.58	9.92	6.26	4.4	5.47		
	NT2RP3004075	4.16	4.16	11.23	12.62	11.88	13.3		
50	NT2RP3004078	2.6	2.6	5.25	4.94	4.19	2.79		
	NT2RP3004083	2.93	2.93	6.23	4.57	6.8	11.37		
	NT2RP3004084	4.65	4.65	20.29	6.18	8.56	5.32		
55	NT2RP3004087	4.2	4.2	7.86	7.14	10.81	9.03		
	NT2RP3004090	4.11	4.11	6.42	9	8.19	8.61	**	+

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	NT2RP3003749	0.67	0.67	2.58	1.55	2.08	1.73		
	NT2RP3003754	3.32	3.32	7.31	4.66	5.87	5.81		
5	NT2RP3003759	1.16	1.16	4.44	2.49	4.41	1.43		
	NT2RP3003764	3.97	3.97	7.08	6.85	7.41	5.06		
	NT2RP3003766	6.93	6.93	7.84	3.3	5.87	3.79	*	-
10	NT2RP3003767	11.19	11.19	16.8	14.83	21.08	16.97		
	NT2RP3003778	3.36	3.36	4.89	4.46	5.55	4.27		
	NT2RP3003779	4.05	4.05	15.26	13.02	8.74	10.52		
	NT2RP3003783	9.25	9.25	21.72	22.42	13.65	18.76		
15	NT2RP3003787	2.15	2.15	4.65	4.41	4.74	6.37		
	NT2RP3003789	5.12	5.12	10.16	11.63	12.19	14.96	*	+
	NT2RP3003795	1.48	1.48	6.48	4.09	2.82	2.24		
	NT2RP3003799	2.67	2.67	5.5	3.08	2.38	1.75		
20	NT2RP3003800	4.36	4.36	5.92	4.14	4.57	6.91		
	NT2RP3003805	8.15	8.15	6.78	8.4	5.48	5.89		
	NT2RP3003809	1.94	1.94	7.2	5.83	5.4	4.82		
25	NT2RP3003819	3.39	3.39	6.07	7.3	5.97	6.35		
	NT2RP3003824	5.69	5.69	10.69	14.08	14.85	13.32	*	+
	NT2RP3003825	9.06	9.06	16.31	12.87	16.88	16.75		
	NT2RP3003828	4.7	4.7	14.38	13.36	15.69	14.55		
30	NT2RP3003831	4.01	4.01	6.38	5.77	6.54	7.23		
	NT2RP3003833	5.12	5.12	7.5	6.44	8.88	6.96		
	NT2RP3003836	6.37	6.37	5.05	5.74	6.47	4.31		
35	NT2RP3003842	2.7	2.7	9.08	6.84	6.51	7.09		
	NT2RP3003843	9.26	9.26	26.77	16.67	12.71	16.2		
	NT2RP3003844	20.38	20.38	46.56	42.84	27.94	44.32		
	NT2RP3003846	4.04	4.04	8.45	8.94	7.18	8.05		
40	NT2RP3003849	2.27	2.27	2.68	2.67	2.73	1.68		
	NT2RP3003862	28.91	28.91	45.63	32	37.58	44.88		
	NT2RP3003870	4.76	4.76	4.81	2.54	2.93	2.05	**	-
45	NT2RP3003874	21.46	21.46	20.88	33.11	47.25	36.44	*	+
	NT2RP3003876	1.62	1.62	8.08	5.45	7.49	6.81		
	NT2RP3003880	1.74	1.74	4.63	5.31	4.66	4.73		
	NT2RP3003889	1.69	1.69	3.04	3.41	3.53	9.53		
50	NT2RP3003891	1.88	1.88	2.98	2.56	3.19	1.37		
	NT2RP3003914	3.1	3.1	7.35	6.88	5.15	7.39		
	NT2RP3003915	5.03	5.03	8.44	9.52	11.35	8.6		
	NT2RP3003918	6.79	6.79	10.39	10.04	13.71	12.42		
55	NT2RP3003920	6.9	6.9	9.13	8.31	10.22	8.96		

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	NT2RP3003576	5.88	5.88	10.15	11.32	8.98	9.22		
	NT2RP3003587	7.39	7.39	12.41	10.01	12.71	12.75		
5	NT2RP3003589	15.33	15.33	22.45	23.89	23.75	26.58		
	NT2RP3003592	7.77	7.77	10.4	8.42	14.48	9.74		
	NT2RP3003593	8.16	8.16	13.62	13.47	13.84	110.49		
10	NT2RP3003614	2.66	2.66	8.18	3.11	4.48	7.09		
	NT2RP3003621	1.64	1.64	3.91	2.1	3.68	2.96		
	NT2RP3003625	1.54	1.54	6.94	3.79	5.09	4.96		
	NT2RP3003627	6.73	6.73	20.05	16.23	13.97	25.71		
15	NT2RP3003636	3.3	3.3	7.74	5.99	3.79	10.4		
	NT2RP3003642	7.12	7.12	12.2	12.85	13.15	15.83		
	NT2RP3003645	2.91	2.91	6.07	2.23	2.42	3.53		
	NT2RP3003648	2.88	2.88	3.71	2.17	2.44	3.13		
20	NT2RP3003649	2.7	2.7	9.28	6.36	5.11	12.04		
	NT2RP3003650	2.65	2.65	4.25	4.38	3.16	4.09		
	NT2RP3003656	1.69	1.69	3.23	1.94	4.12	3		
25	NT2RP3003659	2.76	2.76	4.56	2.14	4.8	4.88		
	NT2RP3003662	31.39	31.39	53.28	34.35	14.68	34.64		
	NT2RP3003664	3.56	3.56	6.5	6.18	5.45	6.55		
	NT2RP3003665	1.89	1.89	4.83	2.07	2.8	4.96		
30	NT2RP3003671	2.88	2.88	4.33	3.03	2.6	4.29		
	NT2RP3003672	4.78	4.78	9.8	10.69	14.73	16.35	*	+
	NT2RP3003673	4.98	4.98	9.42	5.35	3.05	4.12		
35	NT2RP3003679	40.1	40.1	95.75	69.92	23.86	83.88		
	NT2RP3003680	3.13	3.13	5.38	3.96	4.58	5.88		
	NT2RP3003686	2.22	2.22	4.43	2.84	4.85	2.25		
	NT2RP3003689	4.05	4.05	9.69	5.94	5.63	8.27		
40	NT2RP3003697	13.79	13.79	120.74	108.93	77.49	68.74		
	NT2RP3003701	2.7	2.7	5.17	2.58	3.05	2.57		
	NT2RP3003704	2.99	2.99	6.96	7.09	7.61	6.96		
45	NT2RP3003714	1.39	1.39	4.25	1.68	0.89	1.14		
	NT2RP3003716	2.05	2.05	4.23	3	2.29	2.24		
	NT2RP3003721	1.83	1.83	3.27	1.85	3.45	2.18		
	NT2RP3003722	3.45	3.45	8.18	8.08	7.79	5.45		
50	NT2RP3003726	3.5	3.5	4.9	2.77	4.51	2.32		
	NT2RP3003729	4.1	4.1	8.53	4.22	5.44	4.6		
	NT2RP3003731	5.06	5.06	6.98	4.19	3.54	7.45		
55	NT2RP3003740	2.58	2.58	5.08	2.42	2.48	2.94		
	NT2RP3003746	3.63	3.63	8.14	6.7	5.94	6.59		

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	NT2RP3003354	5.09	5.09	13.72	16.29	12.02	13.5		
	NT2RP3003368	3.03	3.03	4.73	4.04	4.08	2.63		
5	NT2RP3003375	4.1	4.1	7.4	7.41	9.67	6.62		
	NT2RP3003377	4.16	4.16	3.98	2.57	3.58	1.65		
	NT2RP3003384	5.77	5.77	4.55	2.83	3.43	2.56	**	-
	NT2RP3003385	4.55	4.55	3.12	1.9	2.36	1.47	*	-
10	NT2RP3003396	3.93	3.93	13.63	16.4	8.38	12.32		
	NT2RP3003403	1.62	1.62	2.54	3.24	4.73	1.92		
	NT2RP3003409	1.18	1.18	2.97	3.3	4.48	3.03		
15	NT2RP3003411	4.59	4.59	15.42	14.11	15.42	10.96		
	NT2RP3003420	3.79	3.79	4.36	3.68	2.13	1.85		
	NT2RP3003425	3.25	3.25	6.71	5.85	7.25	5.49		
	NT2RP3003426	9.11	9.11	16.3	10.88	11.12	17.45		
20	NT2RP3003427	5.95	5.95	10.09	9.15	13.58	8.03		
	NT2RP3003433	2.55	2.55	6.26	8.42	9.57	4.87		
	NT2RP3003437	22.12	22.12	49.85	51.81	44	38.77		
25	NT2RP3003448	1.88	1.88	4.24	3.5	3.83	2.63		
	NT2RP3003455	5.23	5.23	12.16	11.8	9.96	8.44		
	NT2RP3003462	4.96	4.96	10.07	10.76	8.25	7.08		
	NT2RP3003464	3.79	3.79	5.03	3.01	4.76	1.2		
30	NT2RP3003469	4.1	4.1	7.77	6.62	7.56	5.07		
	NT2RP3003473	22.06	22.06	36.6	54.82	69.25	56.46	**	+
	NT2RP3003474	8.26	8.26	23.04	13.23	12.04	13.52		
35	NT2RP3003475	2.84	2.84	4.04	4.55	4.45	3.28		
	NT2RP3003490	2.7	2.7	5.81	4.21	4.43	3.12		
	NT2RP3003491	2.26	2.26	3	2.14	2.75	1.26		
	NT2RP3003493	11.75	11.75	30.77	34.59	28.9	34.45		
40	NT2RP3003500	4.93	4.93	5.26	4.99	7.46	3.65		
	NT2RP3003527	2.73	2.73	3.09	2.42	2.92	1.72		
	NT2RP3003532	2.7	2.7	1.81	2.14	3.33	2.13		
45	NT2RP3003535	3.14	3.14	4.37	1.92	3.4	2.19		
	NT2RP3003536	3.04	3.04	5.95	3.45	5.2	4.97		
	NT2RP3003543	2.61	2.61	6.24	3.21	4.81	3.83		
	NT2RP3003549	1.43	1.43	6.66	2.18	4.07	1.55		
50	NT2RP3003552	1.8	1.8	5.76	0.64	0.84	1.58		
	NT2RP3003555	4.4	4.4	14.14	12.16	17.43	16.23		
	NT2RP3003559	2.81	2.81	6.7	3.88	5.11	5.49		
	NT2RP3003564	3.11	3.11	5.9	2.24	4.6	5.25		
55	NT2RP3003572	2.1	2.1	4.21	1.88	3.02	2.32		

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	NT2RP3003138	2.36	2.36	5.27	5.26	5.7	3.74		
	NT2RP3003139	2.53	2.53	7.8	3.15	5.74	2.7		
5	NT2RP3003145	5.08	5.08	32.56	25.8	29.74	19.72		
	NT2RP3003150	2.03	2.03	5.17	3.56	3.76	1.97		
	NT2RP3003157	2.52	2.52	8.34	6.4	10.1	4.94		
10	NT2RP3003185	1.77	1.77	3.88	1.91	3.34	2.62		
	NT2RP3003193	2.62	2.62	5.75	6.03	4.59	2.65		
	NT2RP3003197	2.38	2.38	3.8	3.11	4.02	2.2		
	NT2RP3003203	11.82	11.82	14.35	16.85	10.17	15.27		
15	NT2RP3003204	3.76	3.76	7.93	4.04	6.17	3.79		
	NT2RP3003210	14.48	14.48	75.3	58.97	84.6	68.66		
	NT2RP3003212	5.15	5.15	9.44	9.21	10.67	7.36		
	NT2RP3003213	4.16	4.16	5.68	5.15	7.02	5.44		
20	NT2RP3003224	1.7	1.7	4.75	2.43	2.11	2.64		
	NT2RP3003226	3.25	3.25	5.68	6.57	5.94	3.63		
	NT2RP3003230	7.79	7.79	11.47	12.39	8.89	6.72		
25	NT2RP3003235	7.61	7.61	10.79	7.77	7.73	6.89		
	NT2RP3003242	12.17	12.17	23.49	26.68	32.03	19.25		
	NT2RP3003251	5.61	5.61	9.47	3.73	4.95	4.08		
	NT2RP3003252	3.95	3.95	5.95	2.19	3.7	2.42		
30	NT2RP3003258	4.92	4.92	7.89	19.94	24.95	15.47	**	+
	NT2RP3003260	4.54	4.54	12.34	13.46	11.52	12.68		
	NT2RP3003264	1.64	1.64	5.99	3.18	4.32	1.86		
35	NT2RP3003273	2.18	2.18	4.93	4.57	3.58	1.72		
	NT2RP3003278	1.33	1.33	4	1.31	5.12	0.63		
	NT2RP3003280	9.85	9.85	23.11	18.18	19.52	18.19		
	NT2RP3003282	5.29	5.29	6.25	3.62	3.97	3.48	**	-
40	NT2RP3003290	6.64	6.64	9.09	4.8	5.38	3.78	*	-
	NT2RP3003301	4.01	4.01	5.73	4.31	4.59	3.23		
	NT2RP3003302	1.45	1.45	2.31	2.91	2.64	1.91		
45	NT2RP3003311	2.45	2.45	6.76	15.72	13.09	11.55	**	+
	NT2RP3003312	1.81	1.81	3.35	3.73	3.87	2.41		
	NT2RP3003313	1.61	1.61	4.2	2.91	5.4	2.87		
	NT2RP3003327	1.62	1.62	6.24	4.81	4.95	3.34		
50	NT2RP3003330	5.13	5.13	8.01	15.68	16.13	12.78	**	+
	NT2RP3003344	3.36	3.36	4.14	2.92	3.74	2.6		
	NT2RP3003346	3.81	3.81	4.83	4.38	4.05	1.24		
	NT2RP3003349	4.04	4.04	6.93	9.96	9.41	9.65	**	+
55	NT2RP3003353	1.95	1.95	3.24	4.06	5.37	2.45		

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	NT2RP3002911	3.05	3.05	3.68	3.09	3.73	2.24		
	NT2RP3002948	4.09	4.09	4.81	2.73	4.44	2.07		
5	NT2RP3002953	3.85	3.85	3.6	2.27	2.52	0.84	*	-
	NT2RP3002955	6.55	6.55	3.78	1.93	2.47	0.86	*	-
	NT2RP3002958	5.85	5.85	11.4	16.5	10.57	16.64		
10	NT2RP3002969	4.28	4.28	8.27	12.91	7.49	7.33		
	NT2RP3002972	3.55	3.55	4.82	4.41	6.18	2.29		
	NT2RP3002978	1.48	1.48	2.99	2.61	3.7	1.49		
	NT2RP3002983	2.89	2.89	4.69	4.46	6.12	4.37		
15	NT2RP3002985	4.23	4.23	17.87	13.64	20.26	11.12		
	NT2RP3002988	3.97	3.97	4.6	4.12	5.64	4.13		
	NT2RP3003000	3.11	3.11	3.46	2.46	3.2	1.51		
20	NT2RP3003008	3.26	3.26	5.87	3.95	4.55	2.96		
	NT2RP3003012	3.43	3.43	6.06	3.9	4.96	2.79		
	NT2RP3003015	1.35	1.35	4.9	1.5	2.5	0.54		
	NT2RP3003018	2.15	2.15	6.09	3.45	7.24	2.59		
25	NT2RP3003028	3.53	3.53	7.23	3.5	5.05	4.01		
	NT2RP3003029	111.75	111.75	149.73	175.13	159.77	181.4	*	+
	NT2RP3003032	7.06	7.06	9.05	11.87	18.84	9.94		
30	NT2RP3003041	2.07	2.07	1.88	1.61	1.41	0.69		
	NT2RP3003044	3.06	3.06	7.45	5.72	6.11	7.57		
	NT2RP3003047	3.09	3.09	5.16	2.4	4.67	2.06		
	NT2RP3003050	5.96	5.96	12.03	6.74	10.3	8.42		
35	NT2RP3003053	7.46	7.46	18	14.42	17.14	13.72		
	NT2RP3003059	1.93	1.93	4.76	2.88	4.41	3.06		
	NT2RP3003061	2.8	2.8	8.59	5.49	5.68	5.16		
40	NT2RP3003068	5.99	5.99	11.77	9.41	8.75	10.04		
	NT2RP3003071	7.22	7.22	10.77	10.39	14.52	10.39		
	NT2RP3003076	2.67	2.67	9.49	6.57	6.57	4.01		
	NT2RP3003078	1.5	1.5	4.12	2.09	4.52	2.43		
45	NT2RP3003081	6.21	6.21	10.54	9.94	9.6	9.97		
	NT2RP3003090	1.49	1.49	5.95	3.28	3.57	3.25		
	NT2RP3003097	2.42	2.42	7.15	2.71	3.72	2.93		
	NT2RP3003098	2.75	2.75	4.22	2.73	3.43	1.73		
50	NT2RP3003101	5.56	5.56	7.24	7.73	10.35	7.18		
	NT2RP3003109	16.11	16.11	27.38	27.36	41.03	21.91		
	NT2RP3003121	3.39	3.39	11.03	4.61	8	2.44		
55	NT2RP3003133	2.09	2.09	5.78	4.93	8.58	3.96		
	NT2RP3003137	3.42	3.42	5.74	6.32	7.59	5.29		

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	NT2RP3002713	4.31	4.31	7.06	2.66	2.19	1.87	*	-
	NT2RP3002721	5.77	5.77	10.06	11.06	16.94	8.96		
5	NT2RP3002722	7.11	7.11	-10.08	7.8	6.45	6.62		
	NT2RP3002723	42.31	42.31	75.85	60.39	46.74	58.76		
	NT2RP3002737	8.35	8.35	18.1	10.97	11.6	9.37		
10	NT2RP3002738	1.9	1.9	6.13	3.09	5.23	3.54		
	NT2RP3002742	14.11	14.11	23.22	30.39	28.27	27.66	*	+
	NT2RP3002744	4.09	4.09	5.24	3.92	4.92	1.71		
	NT2RP3002756	5.8	5.8	5.8	3.19	2.68	1.55	**	-
15	NT2RP3002757	12	12	17.79	19.76	24.24	19.75	*	+
	NT2RP3002758	21.11	21.11	42.35	44.47	63.91	36.38		
	NT2RP3002762	5.07	5.07	8.82	7.21	7.43	7.76		
20	NT2RP3002763	1.62	1.62	4.86	3.76	4.99	2.18		
	NT2RP3002770	1.78	1.78	5.14	3.46	3.7	2.93		
	NT2RP3002771	17.04	17.04	39.53	24.93	40.21	34.4		
	NT2RP3002785	2.42	2.42	5.45	3.36	4.09	2.66		
25	NT2RP3002790	4.65	4.65	4.22	3.16	3.57	2.33	*	-
	NT2RP3002799	4.73	4.73	6.33	3.42	2.7	1.43	*	-
	NT2RP3002801	4.14	4.14	3.59	3.6	3.22	2.49		
30	NT2RP3002802	2.31	2.31	6.3	6.78	5.43	4.4		
	NT2RP3002810	2.98	2.98	5.41	7.44	12.32	13.27	*	+
	NT2RP3002818	1.5	1.5	2.44	2.18	4.16	2.47		
	NT2RP3002821	12.8	12.8	33.14	26.1	35.81	23.02		
35	NT2RP3002823	3.85	3.85	8.98	4.65	5.92	3.87		
	NT2RP3002825	5.47	5.47	13.04	13.47	19.19	7.12		
	NT2RP3002829	5.37	5.37	6.25	4.75	4.89	3.8		
40	NT2RP3002831	4.01	4.01	6.13	9.07	8.77	5.19		
	NT2RP3002836	7.33	7.33	19.42	11.56	16.91	20.66		
	NT2RP3002845	4.17	4.17	6.63	7.87	8.6	8.45	*	+
	NT2RP3002852	3.37	3.37	7.57	7.8	8.72	8.21		
45	NT2RP3002861	3.82	3.82	6.4	7.34	7.35	4.63		
	NT2RP3002869	3.66	3.66	3.26	2.49	1.86	0.49	*	-
	NT2RP3002874	11.25	11.25	21.44	25.33	31.95	25.54	*	+
	NT2RP3002876	6.98	6.98	11.06	12.8	14.93	14.39	*	+
50	NT2RP3002877	4.7	4.7	5.96	3.3	5.24	2.53		
	NT2RP3002887	0.47	0.47	3.42	2.81	3.53	3.91		
	NT2RP3002900	6.46	6.46	19.64	21.86	21.54	22.3		
55	NT2RP3002902	4.01	4.01	10.25	11.72	8.52	8.06		
	NT2RP3002909	2.61	2.61	6.19	6.67	5.38	3.93		

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	NT2RP3002566	3.62	3.62	7.02	4.03	8.51	3.65		
	NT2RP3002571	2.53	2.53	4.85	3.77	5.41	3.74		
5	NT2RP3002572	2.98	2.98	5.28	4.75	4.74	5.21		
	NT2RP3002573	4.31	4.31	11.38	7.06	9.48	7.06		
	NT2RP3002577	1.57	1.57	4.61	2.71	2.32	1.9		
	NT2RP3002579	3.92	3.92	6.41	4.03	7.75	11.16		
10	NT2RP3002582	5.02	5.02	7.17	11.51	14.07	8.45	*	+
	NT2RP3002587	1.9	1.9	3.13	2.68	3.04	1.97		
	NT2RP3002590	3.16	3.16	5.65	9.06	10.39	8.06	**	+
15	NT2RP3002602	3.02	3.02	4.24	3.95	5.85	3.77		
	NT2RP3002603	71.53	71.53	214.41	268.41	257.84	298.26	*	+
	NT2RP3002621	1.95	1.95	3.42	2.13	5.13	1.85		
	NT2RP3002622	2.63	2.63	7.38	3.7	7.36	4.67		
20	NT2RP3002624	2.29	2.29	7.4	4.04	4.9	3.64		
	NT2RP3002628	6.36	6.36	16.17	19.57	22.15	16.21		
	NT2RP3002629	8.96	8.96	13.58	15.4	18.26	15.57	*	+
25	NT2RP3002631	1.95	1.95	1.67	0.91	1.65	1.69		
	NT2RP3002647	4.04	4.04	4.01	5.44	7.16	4.6		
	NT2RP3002649	2.99	2.99	5.99	2.15	5.84	3.23		
	NT2RP3002650	3.32	3.32	11.62	4.98	8.53	6.88		
30	NT2RP3002652	2.27	2.27	6.59	4.91	6.83	4.25		
	NT2RP3002654	3.05	3.05	7.5	5.58	5.4	4.2		
	NT2RP3002657	14.14	14.14	13.87	17.27	26.08	18.87		
35	NT2RP3002659	1.92	1.92	6.01	3.91	5.78	3.47		
	NT2RP3002660	3.09	3.09	4.84	3.77	7.72	3.52		
	NT2RP3002663	2.39	2.39	3.33	2.54	3.13	2.84		
	NT2RP3002664	2.74	2.74	7.28	3.65	2.56	2.13		
40	NT2RP3002667	2.92	2.92	6.59	6.63	5.46	5.35		
	NT2RP3002671	2.37	2.37	5.02	3.91	5.52	4.11		
	NT2RP3002682	6.34	6.34	20.62	14.37	17.64	21.7		
45	NT2RP3002684	4	4	6.34	3.32	6.16	3.18		
	NT2RP3002687	3.25	3.25	6.22	2.7	3.87	6.41		
	NT2RP3002688	3.22	3.22	4.98	2.63	3.91	2.61		
	NT2RP3002698	2.2	2.2	3.99	3.07	4.28	2.38		
50	NT2RP3002701	2.93	2.93	6.73	3.45	3.07	3.6		
	NT2RP3002705	2.17	2.17	8.01	4.36	8.76	4.72		
	NT2RP3002708	3.69	3.69	9.88	5.64	7.34	4.9		
	NT2RP3002711	6.67	6.67	7.85	7.77	7.56	6.69		
55	NT2RP3002712	55.99	55.99	75.28	146.74	168.42	130.64	**	+

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	NT2RP3002352	6.3	6.3	8.01	4.4	6.76	4.31		
	NT2RP3002353	3	3	4.85	4.87	6.18	5.9		
5	NT2RP3002362	5	5	-11.74	15.86	11.03	10.49		
	NT2RP3002363	2.41	2.41	3.67	5.53	6.17	2.32		
	NT2RP3002377	2.61	2.61	5.47	6.8	7.31	4.73		
10	NT2RP3002377	4.47	4.47	7.73	11.4	5.31	7.09		
	NT2RP3002394	5.58	5.58	7.35	7.82	10.17	4.46		
	NT2RP3002397	3.77	3.77	4.81	2.7	3.12	1.68	*	-
	NT2RP3002399	4.61	4.61	7.69	14.65	13.02	16.16	**	+
15	NT2RP3002402	2.84	2.84	6.99	8.94	8.7	6.99		
	NT2RP3002404	2.88	2.88	5.6	3.12	3.73	1.83		
	NT2RP3002410	4.85	4.85	15.65	17.05	14.13	10.65		
20	NT2RP3002411	2.98	2.98	5.68	3.7	5.29	2.85		
	NT2RP3002414	5.62	5.62	9.35	10.28	6.81	7.92		
	NT2RP3002430	5.11	5.11	14.63	18.24	19.29	14.51		
	NT2RP3002448	5.4	5.4	4.6	4.35	5.25	3.62		
25	NT2RP3002454	7.3	7.3	15.31	12.9	12.71	9.32		
	NT2RP3002455	4.62	4.62	12.11	14.31	9.11	13.25		
	NT2RP3002456	3.21	3.21	7.75	7.09	6.57	5.9		
30	NT2RP3002462	2.79	2.79	4.16	4.94	6.17	4.79	*	+
	NT2RP3002469	3.84	3.84	6.38	9.24	6.78	8.07	*	+
	NT2RP3002470	6.7	6.7	14.71	17.86	16.99	13.48		
	NT2RP3002484	4.01	4.01	6.86	6.81	8.01	4.59		
35	NT2RP3002491	3.62	3.62	4.1	2.05	2.51	1.46	**	-
	NT2RP3002494	79.24	79.24	131.02	118.47	163.2	105.2		
	NT2RP3002497	1.07	1.07	1.57	2.37	1.39	1.37		
40	NT2RP3002500	1.23	1.23	1.13	2.72	2.02	2.07	**	+
	NT2RP3002501	5.25	5.25	8.49	8.45	8.11	9.69		
	NT2RP3002512	2.85	2.85	3.97	3.3	2.74	3.57		
	NT2RP3002529	3.94	3.94	7.5	6.59	5.14	5.85		
45	NT2RP3002533	7.95	7.95	10.26	9.79	10.51	8.18		
	NT2RP3002539	4.39	4.39	4.32	5.66	6.61	2.85		
	NT2RP3002540	5.24	5.24	5.5	3.48	4.76	3.65	*	-
	NT2RP3002543	3.44	3.44	7.17	4.93	6.21	5.31		
50	NT2RP3002545	7.34	7.34	7.46	5.17	5.52	6.8	*	-
	NT2RP3002549	3.27	3.27	7.8	4.98	6.11	4.54		
	NT2RP3002552	3.05	3.05	6.04	4.17	5.81	4.06		
55	NT2RP3002558	9.54	9.54	9.39	9.93	4.26	11.27		
	NT2RP3002565	1.94	1.94	4.83	1.73	2.48	1.52		

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	NT2RP3002147	3.8	3.8	10.45	6.06	7.4	6.2	
	NT2RP3002151	5.62	5.62	10.64	6.27	4.43	7.33	
5	NT2RP3002155	1.62	1.62	3.27	2.01	4.98	0.8	
	NT2RP3002156	3.15	3.15	4.82	2.55	4.08	2.42	
	NT2RP3002160	1.57	1.57	3.43	1.36	3.43	1.56	
10	NT2RP3002163	20.86	20.86	55.1	35.13	44.03	32.6	
	NT2RP3002165	4.17	4.17	3.67	6.21	8.31	4.86	
	NT2RP3002166	4.04	4.04	10.53	7.76	8.79	5.58	
	NT2RP3002173	2.24	2.24	5.75	2.95	3.34	3.53	
15	NT2RP3002174	8.41	8.41	15.8	13.21	7.82	14.77	
	NT2RP3002181	1.1	1.1	3.46	1.87	3.51	1.61	
	NT2RP3002185	2.69	2.69	4.51	2.94	4.35	2.61	
20	NT2RP3002193	5.51	5.51	13.38	16.39	15.35	11.36	
	NT2RP3002204	5.66	5.66	12.49	17.04	24.14	18.95	* - +
	NT2RP3002244	4.03	4.03	8.29	5.28	6.11	4.8	
	NT2RP3002248	5.42	5.42	11.1	8.19	11.78	6.52	
25	NT2RP3002253	2.61	2.61	9.3	9.66	11.26	6.18	
	NT2RP3002255	11.07	11.07	26.56	22.78	11.53	20.93	
	NT2RP3002264	3.06	3.06	5.54	5.88	7.37	4.07	
30	NT2RP3002267	1.26	1.26	4.33	3.1	4.65	1.82	
	NT2RP3002273	7.51	7.51	12.98	10.15	13.8	12.11	
	NT2RP3002276	5.22	5.22	7.89	3.08	7.68	3.48	
	NT2RP3002281	6.37	6.37	6.83	7.45	8.46	3.44	
35	NT2RP3002286	3	3	4.79	3.54	4.34	3.88	
	NT2RP3002297	10.62	10.62	29.36	22.26	20.57	23.93	
	NT2RP3002301	5.73	5.73	13.24	9.47	7.55	6.21	
40	NT2RP3002303	3.01	3.01	6.39	5.29	6.65	4.58	
	NT2RP3002304	2.66	2.66	7.17	6.3	7.3	4.91	
	NT2RP3002309	2.3	2.3	7.18	9.26	13	4.39	
	NT2RP3002311	4.54	4.54	6.67	3.17	4.02	1.83	
45	NT2RP3002315	15.27	15.27	20.91	25.82	33.13	21.82	
	NT2RP3002319	2.37	2.37	5.06	3.07	3.51	2.38	
	NT2RP3002324	8.97	8.97	61.42	49.85	51.23	55.4	
	NT2RP3002330	4.74	4.74	8.33	10.31	8.24	8.15	
50	NT2RP3002333	5.13	5.13	14.32	13.14	13.65	8.12	
	NT2RP3002337	2.61	2.61	5.14	4.8	5.57	4.87	
	NT2RP3002342	5.16	5.16	11.56	5.52	7.51	6.43	
55	NT2RP3002343	3.38	3.38	7.29	5.8	7.03	4.13	
	NT2RP3002351	4.32	4.32	4.55	4.38	3.9	2.55	

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	NT2RP3002007	1. 29	1. 29	2. 3	3. 46	4. 05	1. 69		
	NT2RP3002014	1. 38	1. 38	6. 23	6. 04	6. 24	4. 21		
5	NT2RP3002015	3. 61	3. 61	10. 33	14. 17	9. 94	8. 85		
	NT2RP3002033	1. 54	1. 54	5. 03	7. 29	5. 03	3. 65		
	NT2RP3002045	1. 89	1. 89	5. 29	4. 67	4. 36	2. 5		
	NT2RP3002054	5. 26	5. 26	8. 12	6. 27	9. 17	5. 42		
10	NT2RP3002056	5. 67	5. 67	5. 52	4. 24	4. 24	2. 7	*	-
	NT2RP3002057	4. 35	4. 35	3. 5	2. 87	2. 41	0. 81	*	-
	NT2RP3002061	4. 71	4. 71	13. 94	8. 64	8. 9	10. 74		
15	NT2RP3002062	0. 8	0. 8	2. 42	3. 58	3. 26	1. 11		
	NT2RP3002063	5. 61	5. 61	10. 31	9. 29	9. 3	7. 31		
	NT2RP3002064	2. 6	2. 6	3. 37	2. 72	3. 74	2. 52		
	NT2RP3002071	1. 6	1. 6	3. 91	1. 99	3. 29	1. 45		
20	NT2RP3002073	6. 47	6. 47	9. 55	10. 45	11. 13	8. 64		
	NT2RP3002074	4. 2	4. 2	7. 25	6. 33	7. 82	4. 24		
	NT2RP3002075	7. 58	7. 58	11. 93	21. 64	30. 17	18. 15	*	+
25	NT2RP3002077	3. 81	3. 81	5. 95	2. 48	3. 05	2. 78		
	NT2RP3002081	4. 25	4. 25	7. 55	13. 22	12. 62	11. 13	**	+
	NT2RP3002086	3. 86	3. 86	9. 77	5. 59	8. 66	6. 95		
	NT2RP3002094	7. 34	7. 34	10. 28	13. 84	14. 79	11. 67	*	+
30	NT2RP3002096	1. 98	1. 98	4. 53	1. 28	3. 12	1. 73		
	NT2RP3002097	3. 77	3. 77	6. 16	6. 1	8. 34	6. 88		
	NT2RP3002098	1. 61	1. 61	4. 3	1. 04	1. 8	1. 46		
35	NT2RP3002102	2	2	4. 86	3. 11	3. 4	3. 16		
	NT2RP3002106	2. 74	2. 74	4. 98	2. 83	4. 9	2. 51		
	NT2RP3002108	3. 69	3. 69	7. 8	3. 11	3. 39	3. 15		
	NT2RP3002109	12. 49	12. 49	32. 04	31. 61	27. 15	25. 12		
40	NT2RP3002110	36. 38	36. 38	54. 93	55. 24	58. 94	46. 55		
	NT2RP3002113	11. 15	11. 15	13. 99	10. 66	15. 22	11. 44		
	NT2RP3002120	2. 22	2. 22	4. 42	2. 31	4. 13	2. 7		
45	NT2RP3002121	5. 93	5. 93	14. 39	13. 38	14. 39	15. 06		
	NT2RP3002126	34. 03	34. 03	108. 96	121. 18	130. 55	142. 49	*	+
	NT2RP3002128	4. 06	4. 06	8. 23	3. 36	6. 87	3. 92		
	NT2RP3002130	8. 29	8. 29	18. 59	11. 69	10. 7	14. 03		
50	NT2RP3002133	14. 24	14. 24	18. 31	10. 06	8. 3	19. 51		
	NT2RP3002136	10. 32	10. 32	15. 42	12. 49	17. 64	17. 49		
	NT2RP3002140	3. 13	3. 13	6. 35	6. 15	3. 9	4. 16		
55	NT2RP3002142	16. 86	16. 86	50. 85	56. 54	81. 25	62. 65	*	+
	NT2RP3002146	4	4	7. 22	5. 14	9. 31	6. 56		

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	NT2RP3001829	21.63	21.63	43.14	35.64	17.14	24.87		
	NT2RP3001836	7.31	7.31	10.67	15.24	7.26	11.37		
5	NT2RP3001839	18.86	18.86	31.77	31.97	19.23	17.53		
	NT2RP3001844	4.15	4.15	11.37	8.33	9.59	8.54		
	NT2RP3001848	9.61	9.61	52.04	44.52	74.75	43.87		
10	NT2RP3001854	6.41	6.41	11.29	12.86	16.75	13.26	*	+
	NT2RP3001855	2.27	2.27	3.94	1.87	1	1.74		
	NT2RP3001857	3.1	3.1	5.22	5.13	5.6	3.33		
	NT2RP3001858	1.53	1.53	4.45	4.41	6.04	2.97		
15	NT2RP3001861	7.35	7.35	16.34	11.85	14.46	10.42		
	NT2RP3001866	4.35	4.35	9.63	5.52	10.42	7.93		
	NT2RP3001871	4.82	4.82	6.34	5.55	6.38	4.55		
20	NT2RP3001874	6.8	6.8	9.73	7.72	11.19	6.14		
	NT2RP3001878	5.98	5.98	6.35	4.59	6.89	5.02		
	NT2RP3001885	3.21	3.21	5.3	5.65	5.25	4.42		
	NT2RP3001896	1.64	1.64	3.49	3.37	2.13	2.02		
25	NT2RP3001898	9.03	9.03	17.69	14.71	8.69	11.94		
	NT2RP3001899	3.21	3.21	7.33	4.52	6.9	5.81		
	NT2RP3001901	4.58	4.58	9.18	8.19	9.44	9.21		
30	NT2RP3001915	4.84	4.84	11.12	14.09	15.67	14.04	*	+
	NT2RP3001926	2.8	2.8	4.88	2.47	2.45	1.65		
	NT2RP3001929	3.74	3.74	4.06	2.56	3.38	0.86		
	NT2RP3001931	4.63	4.63	5.26	3.9	5.62	2.98		
35	NT2RP3001938	2.27	2.27	5.53	4.93	3.93	4.75		
	NT2RP3001943	3.27	3.27	5.36	5.77	6.5	5.56		
	NT2RP3001944	1.77	1.77	3.72	4.08	5.91	4.34		
	NT2RP3001945	4.25	4.25	12.2	11.86	11.78	6.11		
40	NT2RP3001947	2.94	2.94	5.89	4.06	5.41	3.4		
	NT2RP3001949	4.21	4.21	8.9	10.49	11.08	8.08		
	NT2RP3001952	23.54	23.54	43.64	48.59	88.56	41.86		
45	NT2RP3001954	5.06	5.06	3.68	4.34	4.79	1.85		
	NT2RP3001956	4.97	4.97	9.44	7.76	8.22	6.29		
	NT2RP3001967	3.78	3.78	7.74	6.7	5.37	5.66		
	NT2RP3001969	1.71	1.71	2.91	4.05	4.39	3.62	*	+
50	NT2RP3001976	2.25	2.25	4.67	6.22	6.25	4.84	*	+
	NT2RP3001986	3.55	3.55	3.88	3.43	2.82	2.19		
	NT2RP3001989	3.76	3.76	5.23	2.86	3.58	2.54		
	NT2RP3002002	6.68	6.68	9.47	6.25	8.85	3.86		
55	NT2RP3002004	5.02	5.02	6.23	3.79	5.74	3.55		

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	NT2RP3001646	1.56	1.56	3.7	0.89	2.79	0.95		
	NT2RP3001650	2.06	2.06	5.81	4.86	7.03	2.08		
5	NT2RP3001667	4.66	4.66	11.91	6.93	9.95	5.13		
	NT2RP3001671	2.28	2.28	7.98	7.7	4.69	5.99		
	NT2RP3001672	1.33	1.33	4.55	1.66	1.47	1.72		
10	NT2RP3001676	2.18	2.18	5.02	2.35	3.14	2.24		
	NT2RP3001678	2.86	2.86	9.24	5.12	5.14	6.03		
	NT2RP3001679	6.12	6.12	9.19	6.74	4.73	6.91		
	NT2RP3001682	1.82	1.82	5.09	4.45	6.18	3.35		
15	NT2RP3001685	3.02	3.02	6.74	3.52	6.53	3.01		
	NT2RP3001688	3.01	3.01	9.42	5.46	8.21	6.43		
	NT2RP3001690	3.21	3.21	4.87	2.91	3.54	2.99		
20	NT2RP3001693	5.69	5.69	10.93	16.59	18.34	16.12	**	+
	NT2RP3001696	2.28	2.28	3.63	1.77	3.68	3.39		
	NT2RP3001698	35.35	35.35	79.65	85.09	91.88	105.32		
	NT2RP3001708	4.82	4.82	8.78	6.34	6.95	9.01		
25	NT2RP3001712	8.69	8.69	16.06	10.22	14.19	13		
	NT2RP3001716	1.44	1.44	5.45	2.14	3.42	2.31		
	NT2RP3001724	2.75	2.75	6	4.08	4.54	2.63		
30	NT2RP3001727	11.73	11.73	38.73	39.17	49.36	31.26		
	NT2RP3001729	3.36	3.36	4.7	5.69	6.55	3.06		
	NT2RP3001730	12.54	12.54	26.52	12.53	19.94	16.4		
	NT2RP3001733	1.46	1.46	3.04	2.09	3.7	1.62		
35	NT2RP3001737	3.02	3.02	7.12	4.62	5.49	2.78		
	NT2RP3001738	1.59	1.59	8.22	3.38	6.01	3.03		
	NT2RP3001739	3.26	3.26	5.25	5.63	6.1	2.51		
	NT2RP3001742	2.54	2.54	5.36	3.86	4.55	4.03		
40	NT2RP3001751	3.61	3.61	11.54	9.94	12.82	8.76		
	NT2RP3001752	2.58	2.58	7.01	2.1	3.59	2.76		
	NT2RP3001753	5.73	5.73	9.48	10.83	15.3	13.69	*	+
45	NT2RP3001754	4.63	4.63	9.08	5.86	3.73	5.33		
	NT2RP3001756	4.66	4.66	7.36	9.37	5.75	8.03		
	NT2RP3001764	2.1	2.1	3.76	2.54	4.25	2.49		
	NT2RP3001771	2.63	2.63	3.2	1.52	4.14	1.22		
50	NT2RP3001777	2.59	2.59	5.99	3.25	5.19	3.26		
	NT2RP3001782	3.52	3.52	14.68	6.47	6.63	6.47		
	NT2RP3001792	2.27	2.27	4.35	2.91	4.09	1.35		
55	NT2RP3001799	1.76	1.76	5.18	5.71	6.36	5.68		
	NT2RP3001819	1.36	1.36	4.54	1.7	1.52	2.06		

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	NT2RP3001453	1.84	1.84	3.66	3.5	4.4	2.81		
	NT2RP3001457	3.86	3.86	7.71	6.06	6.93	5.5		
5	NT2RP3001459	2.39	2.39	6.03	2.64	2.78	1.17		
	NT2RP3001463	2.77	2.77	6.74	5.93	5.94	3.98		
	NT2RP3001466	2.87	2.87	3.56	1.19	1.43	0.78	**	-
10	NT2RP3001472	5.74	5.74	4.02	3.7	4.85	4.32		
	NT2RP3001475	3.54	3.54	7.61	6.91	6.65	7.39		
	NT2RP3001479	2.54	2.54	6.66	4.37	5.69	5.16		
	NT2RP3001490	3.18	3.18	9.26	4.4	6.02	5.21		
15	NT2RP3001492	4.36	4.36	7.84	7.59	7.08	5.72		
	NT2RP3001495	4.14	4.14	3.85	2.75	2.92	1.76	*	-
	NT2RP3001497	5.8	5.8	6.32	7.47	9.96	6.8		
20	NT2RP3001501	5.36	5.36	5.52	3.12	4.49	3.43	*	-
	NT2RP3001527	4.89	4.89	6.71	4.9	5.14	3.52		
	NT2RP3001529	1.51	1.51	3.5	4.12	3.95	4.18	*	+
	NT2RP3001538	1.78	1.78	6.2	6.93	7.81	6.23		
25	NT2RP3001539	5.81	5.81	14.5	15.19	14.15	16.47		
	NT2RP3001542	1.52	1.52	5.26	4.23	4.38	2.13		
	NT2RP3001549	4.75	4.75	11.12	14.57	11.37	13.44		
30	NT2RP3001554	3.06	3.06	6.16	6.37	7.5	5.05		
	NT2RP3001560	4.96	4.96	5.73	4.67	6.35	2.36		
	NT2RP3001561	8.85	8.85	20.77	20.38	27.2	17.15		
	NT2RP3001564	1.54	1.54	8.24	6.43	4.53	5.96		
35	NT2RP3001568	2.1	2.1	7.68	11.84	10.29	8.49	*	+
	NT2RP3001575	3.94	3.94	7.24	6.39	6.97	6.16		
	NT2RP3001580	1.78	1.78	4.49	4.35	3.8	3.11		
40	NT2RP3001587	4.38	4.38	8.74	10.75	10.04	7.77		
	NT2RP3001589	3.17	3.17	8.21	5.6	7.79	4.36		
	NT2RP3001592	4.52	4.52	21.6	19	32.62	14.54		
	NT2RP3001607	3.42	3.42	1.86	1.59	2.8	1		
45	NT2RP3001608	1.05	1.05	3.59	2.41	1.73	2.31		
	NT2RP3001613	3.08	3.08	2.77	3.89	2.91	3.99		
	NT2RP3001619	4.31	4.31	8.15	7.69	6.45	7.62		
	NT2RP3001621	1.18	1.18	2.69	2.39	2.28	2.02		
50	NT2RP3001629	2.58	2.58	3.28	2.68	2.41	1.7		
	NT2RP3001630	3.39	3.39	4.56	1.67	2.02	1.17	**	-
	NT2RP3001631	9.01	9.01	14.34	18.65	21.16	15.24	*	+
55	NT2RP3001634	4	4	5.29	4.51	6.89	5.11		
	NT2RP3001642	3.71	3.71	7.45	5.77	4.41	5.09		

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	NT2RP3001301	2.91	2.91	6.59	3.96	4.58	3.9		
	NT2RP3001307	1.76	1.76	7.67	2.07	2.81	2.06		
5	NT2RP3001310	11.55	11.55	-17.04	25.54	26.07	28.13	**	+
	NT2RP3001318	2.11	2.11	3.4	2.49	3.37	2.37		
	NT2RP3001322	3.58	3.58	5.23	2.62	3.84	5.48		
	NT2RP3001325	2.7	2.7	8.39	5.82	6.82	5.58		
10	NT2RP3001338	2.67	2.67	6.19	4.1	4.21	3.5		
	NT2RP3001339	2.53	2.53	5.64	3.08	4.89	2.91		
	NT2RP3001340	2.9	2.9	8.42	6.36	7.07	5.79		
15	NT2RP3001341	2.26	2.26	6.97	5.1	5.62	4.73		
	NT2RP3001354	3.22	3.22	9.77	4.28	6.93	9.35		
	NT2RP3001355	1.9	1.9	5.41	2.65	3.82	2.74		
	NT2RP3001356	2	2	5.34	2.59	3.2	3.55		
20	NT2RP3001359	1.09	1.09	4.05	1.63	2.5	1.75		
	NT2RP3001364	2.34	2.34	5.31	3.26	6.67	2.67		
	NT2RP3001373	1.12	1.12	3.22	2.1	3.74	1.71		
25	NT2RP3001374	1.9	1.9	4.17	3.18	3.92	3.1		
	NT2RP3001383	3.84	3.84	8.96	3.92	6.65	3.85		
	NT2RP3001384	4.11	4.11	9.47	3.54	4.46	2.41		
	NT2RP3001388	3.98	3.98	8.79	9.48	10.99	9.4		
30	NT2RP3001392	4.61	4.61	6.19	3.91	6.14	3.23		
	NT2RP3001396	1.7	1.7	6.39	4.04	4.66	4.53		
	NT2RP3001398	2.51	2.51	6.55	3.85	7.05	2.94		
35	NT2RP3001399	4.91	4.91	20.67	15.86	16.12	12.44		
	NT2RP3001402	6.46	6.46	36.36	33.37	41.61	39.66		
	NT2RP3001407	6.96	6.96	19.16	13.69	17.65	12.35		
	NT2RP3001416	7.92	7.92	15.88	13.02	18.3	14.72		
40	NT2RP3001420	5.33	5.33	6.4	3.27	3.64	1.8	*	-
	NT2RP3001425	3.73	3.73	4.92	4.74	5.67	3.15		
	NT2RP3001426	2.39	2.39	6.08	5.45	4.45	5.11		
45	NT2RP3001427	1.82	1.82	5.61	3.46	2.89	3.59		
	NT2RP3001428	2.42	2.42	6.29	5.69	4.81	3.77		
	NT2RP3001429	3.08	3.08	5.91	4.15	7.37	4.73		
	NT2RP3001432	2.14	2.14	6.61	3.72	4.44	3.58		
50	NT2RP3001439	4.14	4.14	6.39	5.87	7.27	4.41		
	NT2RP3001441	6.45	6.45	12.63	11.13	14.61	11.2		
	NT2RP3001446	4.99	4.99	4.99	4.64	5.22	4.39		
55	NT2RP3001447	2.72	2.72	5.21	6.64	5.14	6.33		
	NT2RP3001449	3.95	3.95	11.85	16.9	14.57	13.16	*	+

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	NT2RP3001120	5.02	5.02	8.24	8.85	7.87	6.71		
	NT2RP3001126	6.16	6.16	12.34	17.84	19.66	17.49	**	+
5	NT2RP3001127	6.93	6.93	6.76	4.79	7.63	6.36		
	NT2RP3001133	3.95	3.95	4.95	3.95	4.16	3.62		
	NT2RP3001140	1.46	1.46	2.43	3.21	2.38	6.71		
10	NT2RP3001147	3.16	3.16	6.96	16.08	14.49	13.84	**	+
	NT2RP3001150	1.99	1.99	4.32	4.06	5.68	3.76		
	NT2RP3001152	1.7	1.7	3.29	3.15	3.62	2.51		
	NT2RP3001155	2.95	2.95	4.35	3.68	4.35	3.58		
15	NT2RP3001156	4.38	4.38	6.57	2.91	5.72	5.67		
	NT2RP3001159	5.38	5.38	10.5	7.87	10.86	7.25		
	NT2RP3001170	7.38	7.38	5.96	6.12	8.01	4.56		
20	NT2RP3001176	3.49	3.49	10.75	6.27	8.23	9.49		
	NT2RP3001195	2.35	2.35	4.81	6.79	5.79	6.22	*	+
	NT2RP3001209	3.47	3.47	5.98	5.96	4.64	5.22		
	NT2RP3001214	1.63	1.63	4.91	3.44	3.87	3.81		
25	NT2RP3001216	3.58	3.58	6.38	6.25	4.33	3.6		
	NT2RP3001221	3.33	3.33	4.27	3.07	3.06	1.79		
	NT2RP3001226	5.96	5.96	29.04	21.93	31.45	17.76		
30	NT2RP3001230	3.17	3.17	2.41	3.09	3.14	1.56		
	NT2RP3001232	1.8	1.8	4.72	2.36	3.7	2.85		
	NT2RP3001236	1.68	1.68	4.3	1.7	3.26	1.47		
	NT2RP3001239	1.58	1.58	5.21	2.81	4.31	2.01		
35	NT2RP3001240	12.83	12.83	22.18	23.01	24.3	14.46		
	NT2RP3001245	3.53	3.53	9.88	4.08	6.36	3.39		
	NT2RP3001253	2.79	2.79	4.87	3.34	4.53	5.21		
40	NT2RP3001259	6.62	6.62	11.97	12.33	15.62	11.83		
	NT2RP3001260	3.74	3.74	5.15	3.45	5.44	3.97		
	NT2RP3001264	2.2	2.2	10.29	5.99	6.92	6.38		
	NT2RP3001268	2.25	2.25	7.18	4.93	4.72	4.35		
45	NT2RP3001271	7.06	7.06	16.29	13.07	12.27	14.24		
	NT2RP3001272	3.73	3.73	12.45	9.43	11.09	10.15		
	NT2RP3001274	6.08	6.08	8.09	6.72	6.35	5.11		
	NT2RP3001275	9.78	9.78	11.58	21.56	26.84	22.59	**	+
50	NT2RP3001280	3.39	3.39	5.5	3.58	5.24	4.18		
	NT2RP3001281	3.15	3.15	3.89	3.08	4.48	5.14		
	NT2RP3001288	49.31	49.31	103.24	124.07	142.92	164.41	*	+
	NT2RP3001297	6.39	6.39	42.01	37.04	42.75	41.14		
55	NT2RP3001300	5.23	5.23	15.92	16.78	17.41	17.76		

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	NT2RP3000904	3.83	3.83	8.67	3.05	4.87	3.16		
	NT2RP3000907	5.66	5.66	10.03	8.94	10.67	10.14		
5	NT2RP3000913	6.04	6.04	15.01	17.87	25.57	17.37	*	+
	NT2RP3000917	7.64	7.64	16.58	7.66	6.56	13.51		
	NT2RP3000919	1.99	1.99	5.15	3.5	4.3	2.68		
	NT2RP3000921	2.26	2.26	7.67	4.88	4.07	6.51		
10	NT2RP3000942	2.66	2.66	3.89	2.68	4.12	2.63		
	NT2RP3000968	70.24	70.24	87.55	105.89	110.05	115.38	**	+
	NT2RP3000974	5.36	5.36	9.06	3.21	4.48	2.64		
15	NT2RP3000980	5.77	5.77	5.77	2.09	4.14	2.26	*	-
	NT2RP3000984	3.17	3.17	7.65	6.33	6.68	4.58		
	NT2RP3000994	2.09	2.09	4.88	2.4	3.14	3.37		
	NT2RP3001001	1.46	1.46	3.45	3.75	4.14	1.31		
20	NT2RP3001004	3.37	3.37	6.52	3.51	5.63	5.01		
	NT2RP3001007	4.46	4.46	9.87	10.02	10.62	6.81		
	NT2RP3001012	2.78	2.78	5.4	4.99	6.83	3.65		
25	NT2RP3001042	4.74	4.74	5.52	2.99	5.18	1.38		
	NT2RP3001044	6.26	6.26	7.12	7.16	6.76	4.92		
	NT2RP3001048	2.52	2.52	3.01	3.5	3.93	2.42		
	NT2RP3001050	1.79	1.79	4.99	4.68	5.94	4.7		
30	NT2RP3001055	6.55	6.55	15.6	16.48	12.44	20.49		
	NT2RP3001057	2.79	2.79	10.84	5.57	6.05	5.67		
	NT2RP3001061	3.18	3.18	6.57	5.03	7.85	5.42		
35	NT2RP3001069	6.03	6.03	14.95	18.49	17.53	15.08		
	NT2RP3001074	4.2	4.2	7.22	8.72	10	6.64		
	NT2RP3001078	5.11	5.11	7.29	7.51	8.72	5.18		
	NT2RP3001081	4	4	5.72	4.65	5.19	3.52		
40	NT2RP3001084	2.7	2.7	7.92	6.85	6.71	6.23		
	NT2RP3001095	1.57	1.57	3.88	3.69	3.68	3.42		
	NT2RP3001096	2.52	2.52	7.33	16.78	8.08	18.7	*	+
45	NT2RP3001097	3.65	3.65	4.28	6.42	8.11	8.5	**	+
	NT2RP3001107	3.69	3.69	4.79	3.77	4.03	2.37		
	NT2RP3001109	3.2	3.2	5.5	6.01	9.56	7.26	*	+
	NT2RP3001111	4.58	4.58	4.19	3.41	3.51	2.29	*	-
50	NT2RP3001112	12.61	12.61	18.48	25.73	29.85	24.61	**	+
	NT2RP3001113	1.21	1.21	2.59	2.47	3.24	2.19		
	NT2RP3001115	1.51	1.51	3.32	2.57	3.77	2.19		
	NT2RP3001116	1.01	1.01	2.66	2.55	4.4	2.91		
55	NT2RP3001119	3.69	3.69	6.75	9.07	6.67	5.44		

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	NT2RP3000721	5.63	5.63	24.61	23.43	39.76	21.55		
	NT2RP3000728	3.33	3.33	2.57	1.4	1.64	1.05	**	-
5	NT2RP3000730	2.06	2.06	-5.04	2.76	4.23	1.86		
	NT2RP3000733	2.87	2.87	6.32	3.48	4.47	4.25		
	NT2RP3000735	1.74	1.74	4.22	1.81	2.22	1.26		
10	NT2RP3000736	2.71	2.71	6.35	3.29	5.05	3.65		
	NT2RP3000739	13.76	13.76	12.16	18.05	9.37	20.19		
	NT2RP3000742	3.89	3.89	10.06	4.54	4.97	4.43		
	NT2RP3000753	2.29	2.29	3.9	2.17	2.65	6.3		
15	NT2RP3000759	9.07	9.07	15.99	11.11	17.14	23.05		
	NT2RP3000789	1.58	1.58	5.76	4.89	4.23	3.69		
	NT2RP3000815	1.91	1.91	5.92	4.49	5.57	3.08		
20	NT2RP3000818	4.35	4.35	11.29	6.64	10.49	8.27		
	NT2RP3000820	9.01	9.01	18.49	18.58	20.1	16.9		
	NT2RP3000821	2.13	2.13	4.83	3.28	5.19	2.02		
	NT2RP3000825	1.87	1.87	4.94	1.92	1.47	2.27		
25	NT2RP3000826	4.04	4.04	13.59	10.86	13.8	12.94		
	NT2RP3000836	5.33	5.33	11.61	11.55	14.11	13.3		
	NT2RP3000838	319.2	319.2	741.74	710.2	743.55	1049.86		
30	NT2RP3000839	2.35	2.35	6.67	4.53	6.38	4.36		
	NT2RP3000841	2.17	2.17	4.32	3.79	5.55	4.72		
	NT2RP3000845	3.96	3.96	8.89	5.76	6.71	7.85		
	NT2RP3000847	3.7	3.7	7.94	4.48	5.94	5.28		
35	NT2RP3000848	2.84	2.84	8.34	5.36	6.81	6.3		
	NT2RP3000850	5.67	5.67	7.04	6.58	11.29	7.47		
	NT2RP3000852	3.27	3.27	3.17	4.02	5.23	5.8	*	+
40	NT2RP3000859	2.76	2.76	7.12	4.46	7.11	8.43		
	NT2RP3000861	2.58	2.58	10.51	6.13	10.36	6.43		
	NT2RP3000862	15.29	15.29	24.16	16.36	9.81	23.13		
	NT2RP3000865	1.58	1.58	4.26	2.54	4.21	1.83		
45	NT2RP3000866	2.08	2.08	5.03	2.37	3.59	5.22		
	NT2RP3000868	2.2	2.2	7.09	3.04	3.84	2.28		
	NT2RP3000869	3.54	3.54	11.36	9.61	15.76	7.9		
50	NT2RP3000871	1.75	1.75	3.79	1.81	3.24	1.94		
	NT2RP3000875	0.99	0.99	4.25	2.57	2.71	3.64		
	NT2RP3000895	2.54	2.54	5.56	2.84	3.55	4.93		
	NT2RP3000900	6.01	6.01	11.86	11.3	7.7	14.58		
55	NT2RP3000901	3.67	3.67	7.03	4.11	6.39	5.3		
	NT2RP3000903	3.76	3.76	7.87	3.12	5.92	3.93		

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	NT2RP3000542	6.23	6.23	8.1	7.21	7.3	6.39		
	NT2RP3000554	8.81	8.81	15.22	13.78	10.56	14.95		
5	NT2RP3000561	1.21	1.21	3.51	3.11	2.76	2.25		
	NT2RP3000562	1.84	1.84	3.5	3.7	3.87	3.23		
	NT2RP3000578	1.56	1.56	2.54	2.54	3.37	2.36		
	NT2RP3000582	1.26	1.26	4.66	2.24	2.52	0.41		
10	NT2RP3000584	2.82	2.82	6.52	3.2	2.5	2.02		
	NT2RP3000586	4.08	4.08	4.59	3.28	3.9	2.87		
	NT2RP3000590	5.69	5.69	4.61	3.78	4.35	2.57		
15	NT2RP3000592	1.8	1.8	2.99	2.97	2.75	3.15		
	NT2RP3000596	2.27	2.27	4.89	4.5	3.33	3.03		
	NT2RP3000599	1.67	1.67	3.07	3.88	4.98	3.82	*	+
	NT2RP3000603	6.09	6.09	39.25	40.43	44.88	35.89		
20	NT2RP3000605	2.84	2.84	6.66	4.56	4.23	2.56		-
	NT2RP3000607	5.35	5.35	7.59	5.74	8.46	7.55		
	NT2RP3000616	3.26	3.26	5.45	2.56	2.38	1.21		
25	NT2RP3000621	5.18	5.18	8.48	10.28	10.29	6.01		
	NT2RP3000622	2.36	2.36	8.76	5.85	6.21	4.72		
	NT2RP3000624	1.53	1.53	3.19	3.97	3.06	2.78		
	NT2RP3000628	2.44	2.44	8.04	10.27	7.85	5.58		
30	NT2RP3000631	4.71	4.71	14.95	22.82	16.45	14.2		
	NT2RP3000632	2.35	2.35	5.5	7.78	8.91	5.91	*	+
	NT2RP3000638	6.95	6.95	17.93	11.8	11.6	9.97		
35	NT2RP3000644	25.72	25.72	48.41	57.98	72.01	52.49	*	+
	NT2RP3000645	5.85	5.85	10.48	9.84	12.55	8.43		
	NT2RP3000652	3.39	3.39	5.34	6.22	5.9	7.74	*	+
	NT2RP3000658	2.26	2.26	5.01	6.16	4.24	4.86		
40	NT2RP3000660	2.34	2.34	6.25	6.98	6.91	5.14		
	NT2RP3000661	1.98	1.98	4.49	4.06	3.87	3.1		
	NT2RP3000665	4.79	4.79	12.26	11.83	11.92	7		
45	NT2RP3000676	4.46	4.46	7.55	6.65	7.81	5.42		
	NT2RP3000677	2.87	2.87	4.13	2.44	3.07	1.54		
	NT2RP3000681	19.85	19.85	30.12	32.94	41.51	34.34	*	+
	NT2RP3000683	2.68	2.68	9.67	6.69	7.09	6.69		
50	NT2RP3000685	1.7	1.7	2.5	3.63	2.36	3.44		
	NT2RP3000690	2.77	2.77	3.29	3.82	3.75	2.72		
	NT2RP3000698	10	10	22.49	25.66	17.08	27.43		
55	NT2RP3000708	3.45	3.45	5.5	8.17	9.22	8.56	**	+
	NT2RP3000719	2.83	2.83	2.83	1.16	1.7	1.91	**	-

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	NT2RP3000348	444.59	444.59	802.63	824.62	1016.01	909.68		
	NT2RP3000350	4.25	4.25	10.34	4.57	9.28	6.4		
5	NT2RP3000359	9.53	9.53	24.44	8.54	11.36	16.62		
	NT2RP3000361	7.5	7.5	11.12	7.89	7.81	8.95		
	NT2RP3000366	7.38	7.38	14.27	9.52	11.84	16.13		
	NT2RP3000378	2.67	2.67	5.75	3.92	4.78	2.47		
10	NT2RP3000384	5.42	5.42	10.88	9.52	13.1	9.28		
	NT2RP3000389	12.54	12.54	21.49	23.95	35.02	27.32	*	+
	NT2RP3000393	3.74	3.74	6.16	5.03	4.53	4.77		
15	NT2RP3000395	110.27	110.27	212	108.33	38.18	148.45		
	NT2RP3000397	2.83	2.83	5.28	2.51	5.26	3.31		
	NT2RP3000398	3.39	3.39	10.12	11.46	11.18	12.26		
	NT2RP3000403	3.22	3.22	9.39	10.1	8.2	8.44		
20	NT2RP3000418	3.4	3.4	10.22	7.12	11.08	13.42		-
	NT2RP3000424	2.86	2.86	9.43	6.25	9.52	6.86		
	NT2RP3000427	4.65	4.65	9.05	11.55	13.43	12.35	*	+
25	NT2RP3000431	2.05	2.05	4.93	3.43	3.26	3.93		
	NT2RP3000433	2.63	2.63	8.65	5.65	7.09	6.65		
	NT2RP3000436	11.39	11.39	20.93	18.76	9.35	18.86		
	NT2RP3000439	1.4	1.4	3.61	2.54	3.56	2		
30	NT2RP3000441	3.88	3.88	7.4	7.56	7.92	6.39		
	NT2RP3000444	3.31	3.31	7.29	2.36	3.25	2.2		
	NT2RP3000448	4.45	4.45	10.15	4.05	6.54	3.93		
35	NT2RP3000449	2.84	2.84	4.59	3.1	3.94	2.93		
	NT2RP3000451	1.76	1.76	5.12	3.7	5	2.96		
	NT2RP3000456	1.69	1.69	5.48	4.23	6.67	4.21		
	NT2RP3000460	18.87	18.87	36.67	24.52	25.24	26.25		
40	NT2RP3000471	3.14	3.14	6.49	2.74	4.98	5.84		
	NT2RP3000477	19.96	19.96	23.67	28.98	17.78	32.78		
	NT2RP3000478	5.86	5.86	8.95	5.21	8.98	2.6		
45	NT2RP3000481	5.48	5.48	5.76	2.76	3.61	1.52	**	-
	NT2RP3000484	3.51	3.51	4.26	2.32	2.55	1.76	*	-
	NT2RP3000487	1.77	1.77	7.4	5.07	4.03	4.97		
	NT2RP3000512	3.29	3.29	17.7	15.17	15.9	14.52		
50	NT2RP3000523	13.05	13.05	30.74	31.75	27.83	34.4		
	NT2RP3000526	3.07	3.07	7.38	5.18	6.31	4.64		
	NT2RP3000527	2.83	2.83	6.5	3.76	7.25	5.03		
	NT2RP3000531	2.9	2.9	7.71	5.11	5.51	4.69		
55	NT2RP3000532	5.74	5.74	5.6	5.75	8.39	4.26		

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	NT2RP3000172	0.86	0.86	1.53	1.66	1.2	1.46		
	NT2RP3000186	4.32	4.32	10.6	19.18	15.43	15.82	*	+
5	NT2RP3000197	1.22	1.22	-3.66	4.03	4.29	3.39		
	NT2RP3000201	2.4	2.4	7.2	10.49	8.4	7.88		
	NT2RP3000204	2.16	2.16	4.44	3.88	4.1	4.25		
10	NT2RP3000207	2.87	2.87	4.71	3	2.6	2.45		
	NT2RP3000216	5.38	5.38	10.1	5.87	9.5	5.73		
	NT2RP3000220	5.14	5.14	5.66	3.68	5.69	2.92		
	NT2RP3000221	2.18	2.18	5.45	6.26	6.63	5.93		
15	NT2RP3000232	2.7	2.7	8.01	7.1	5.52	5.92		
	NT2RP3000233	1.55	1.55	6.01	6.9	5.91	4.06		
	NT2RP3000234	3.23	3.23	9.09	12.89	10.4	11.41	*	+
20	NT2RP3000235	1.57	1.57	3.3	2.35	2.92	1.38		
	NT2RP3000239	4.61	4.61	11.11	9.51	9.71	14.92		
	NT2RP3000247	3.25	3.25	5.82	2.92	4.04	1.96		
	NT2RP3000251	6.11	6.11	6.52	5.22	5.82	3.25		
25	NT2RP3000252	3.73	3.73	7.99	7.61	8.53	8.4		
	NT2RP3000255	2.18	2.18	2.96	3.26	3.13	1.97		
	NT2RP3000262	6.72	6.72	9.43	11.67	7.95	9.13		
30	NT2RP3000266	6.47	6.47	15.5	13.38	10.83	12.64		
	NT2RP3000267	2.71	2.71	4.04	2.9	2.64	3.03		
	NT2RP3000271	4.38	4.38	5.57	5.11	4.84	3.72		
	NT2RP3000278	7.84	7.84	56.85	48.55	82.07	42.57		
35	NT2RP3000281	4.94	4.94	10.72	8.19	8.22	7.27		
	NT2RP3000292	5.63	5.63	14.1	9.17	6.77	6.93		
	NT2RP3000299	2.31	2.31	4.92	3.73	4.89	4.98		
	NT2RP3000304	2.15	2.15	3.48	2.85	3.36	1.64		
40	NT2RP3000310	7.24	7.24	24.22	18.94	23.07	19.88		
	NT2RP3000312	2.99	2.99	8.16	3.31	5.25	3.87		
	NT2RP3000320	7.06	7.06	6.17	5.25	4.74	4.74	**	-
45	NT2RP3000322	11.05	11.05	18.76	32.59	45.13	46.95	**	+
	NT2RP3000324	6.91	6.91	46.42	36.64	43.53	39.68		
	NT2RP3000326	1.95	1.95	6.17	4.02	5.75	3.53		
	NT2RP3000329	2.5	2.5	5.96	4.97	8.84	5.9		
50	NT2RP3000330	4.1	4.1	6.18	4.62	5.53	6.12		
	NT2RP3000333	3.23	3.23	7.45	4.36	5.28	4.52		
	NT2RP3000341	8.8	8.8	12.85	14.81	18.59	14.41	*	+
55	NT2RP3000344	2.73	2.73	3.75	2.69	3.54	2.29		
	NT2RP3000345	3.09	3.09	3.57	1.65	1.97	2.66	*	-

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	NT2RP3000014	3.29	3.29	7.66	4.22	3.06	4.95		
	NT2RP3000016	3.42	3.42	7	5.29	6.56	6.11		
5	NT2RP3000022	1.71	1.71	3.93	1.72	4.68	0.59		
	NT2RP3000024	3.74	3.74	7.03	4.31	4.92	4.06		
	NT2RP3000031	4.66	4.66	8.66	4	6.75	4.86		
	NT2RP3000034	3.76	3.76	6.24	4.44	7.13	3.23		
10	NT2RP3000037	2.76	2.76	6.5	9.41	13.44	11.06	*	+
	NT2RP3000040	2.04	2.04	5.96	3.21	3.46	3.56		
	NT2RP3000041	2.15	2.15	7.35	3.71	3.01	3.16		
15	NT2RP3000046	1.95	1.95	4.42	3.67	7.11	3.84		
	NT2RP3000047	3.25	3.25	5.55	5.85	6.2	5.94		
	NT2RP3000049	2.54	2.54	6.94	5.26	7.78	3.58		
	NT2RP3000050	4.99	4.99	9.03	3.76	8.5	6.22		
20	NT2RP3000051	5.99	5.99	10.69	8.51	11.19	9.72		
	NT2RP3000054	4.31	4.31	6.5	4.38	5.35	3.22		
	NT2RP3000055	1.98	1.98	4.76	3.81	2.67	3.96		
25	NT2RP3000056	2.87	2.87	7.09	5.59	3.32	3.91		
	NT2RP3000059	2.54	2.54	5.1	1.89	4.07	1.6		
	NT2RP3000063	2.18	2.18	5.51	3.34	5.19	2.27		
	NT2RP3000068	3.76	3.76	24.22	25.83	37.88	23.13		
30	NT2RP3000069	17.44	17.44	20.58	22	28.87	18.2		
	NT2RP3000072	5.9	5.9	6.18	4.96	5.39	4.19	*	-
	NT2RP3000080	4.38	4.38	6.72	3.78	5.28	3.93		
35	NT2RP3000085	1.9	1.9	4.84	5.13	4.66	5.5		
	NT2RP3000087	3.77	3.77	9.1	6.22	5.61	6		
	NT2RP3000092	1.92	1.92	3.6	2.72	3.2	2.52		
	NT2RP3000109	1.74	1.74	5.05	5.63	7.94	4.24		
40	NT2RP3000119	4.66	4.66	14.27	11.29	13.7	14.28		
	NT2RP3000125	3.02	3.02	5.56	3.42	4.53	2		
	NT2RP3000131	7.84	7.84	14.37	16.23	19.96	12.93		
45	NT2RP3000134	5.96	5.96	9.01	6.61	7.25	6.46		
	NT2RP3000137	3.88	3.88	6.48	5.58	6.3	6.11		
	NT2RP3000142	2.87	2.87	7.77	7.28	5.03	5.31		
	NT2RP3000148	1.84	1.84	6.28	4.9	5.04	5.34		
50	NT2RP3000149	2.51	2.51	6.97	6.14	7.77	8.24		
	NT2RP3000163	2.16	2.16	6.17	3.27	3.9	2.5		
	NT2RP3000168	5.53	5.53	14.55	12.8	11.65	11.73		
	NT2RP3000169	3.74	3.74	6.01	6.03	8.47	5.72		
55	NT2RP3000171	10.86	10.86	16.71	28.33	38.98	25.93	*	+

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	NT2RP2006334	2. 24	2. 24	4. 02	4. 57	2. 72	3. 54		
	NT2RP2006338	2. 4	2. 4	5. 26	4. 73	5. 11	4. 04		
5	NT2RP2006339	2. 24	2. 24	2. 94	2. 47	1. 93	2. 06		
	NT2RP2006355	3. 61	3. 61	4. 59	3. 14	3. 39	2. 22		
	NT2RP2006365	3. 3	3. 3	4. 44	2. 42	2. 6	1. 3	*	-
10	NT2RP2006374	16. 34	16. 34	111. 62	108. 73	174. 7	73. 65		
	NT2RP2006393	4. 93	4. 93	7. 68	7. 38	8	6. 95		
	NT2RP2006394	8. 59	8. 59	17. 91	11. 3	11. 18	15. 38		
	NT2RP2006400	2. 25	2. 25	4. 51	2. 08	3. 58	1. 95		
15	NT2RP2006411	27. 71	27. 71	42. 11	23. 61	17. 25	37. 31		
	NT2RP2006429	2. 22	2. 22	7. 3	2. 82	5. 3	2. 21		
	NT2RP2006435	1. 46	1. 46	5. 29	1. 76	2. 65	1. 98		
20	NT2RP2006436	2. 33	2. 33	6. 43	4. 33	5. 28	3. 75		
	NT2RP2006441	4. 69	4. 69	8. 19	7. 76	8. 89	8. 37		
	NT2RP2006447	2. 41	2. 41	4. 78	3. 18	2. 63	3. 87		
	NT2RP2006454	2. 58	2. 58	5. 38	4. 39	3. 37	4. 03		
25	NT2RP2006455	3. 79	3. 79	7. 14	2. 91	4. 62	9. 23		
	NT2RP2006456	1. 96	1. 96	5. 99	2. 51	4. 49	3. 17		
	NT2RP2006464	5. 44	5. 44	8. 28	4. 47	8. 85	7. 9		
30	NT2RP2006467	4. 17	4. 17	10	8. 56	12. 47	12. 58		
	NT2RP2006472	5. 05	5. 05	6. 84	7. 24	6. 92	7. 37		
	NT2RP2006474	4. 69	4. 69	16. 3	18. 19	32. 31	21. 3		
	NT2RP2006475	2. 5	2. 5	9. 54	6. 14	6. 86	7. 66		
35	NT2RP2006476	5. 34	5. 34	14. 94	7. 62	13. 82	17. 24		
	NT2RP2006501	2. 44	2. 44	7. 28	4. 6	7. 45	7. 74		
	NT2RP2006512	10. 25	10. 25	19. 79	16. 72	7. 89	29. 01		
40	NT2RP2006526	2. 09	2. 09	5. 19	2. 24	2. 78	2. 31		
	NT2RP2006527	3. 61	3. 61	7. 05	4. 56	6. 14	6. 46		
	NT2RP2006534	2. 24	2. 24	4. 49	2. 08	2. 95	2. 73		
	NT2RP2006537	6. 08	6. 08	15. 7	11. 72	17. 73	12. 82		
45	NT2RP2006543	7. 83	7. 83	14. 8	6. 52	5. 4	6. 88		
	NT2RP2006554	1. 33	1. 33	3. 71	1. 79	3. 76	2. 2		
	NT2RP2006565	3. 78	3. 78	8. 91	5. 79	8. 42	7. 55		
	NT2RP2006571	1. 38	1. 38	3. 88	2. 77	4. 01	2. 29		
50	NT2RP2006573	2. 1	2. 1	4. 02	3. 05	3. 6	2. 41		
	NT2RP2006598	2. 25	2. 25	7. 04	4. 34	6. 56	4. 78		
	NT2RP2006601	24. 92	24. 92	35. 13	38. 45	45. 47	31. 69		
55	NT2RP3000002	5. 04	5. 04	6. 09	4. 7	5. 04	8. 18		
	NT2RP3000011	1. 82	1. 82	5. 9	2. 59	1. 85	2. 22		

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	NT2RP2006064	4.16	4.16	7.73	5.86	6.81	9.08		
	NT2RP2006068	2.76	2.76	6.75	6.8	7.81	5.81		
5	NT2RP2006069	1.46	1.46	-4.94	3.56	3.95	3.3		
	NT2RP2006071	8.37	8.37	7.8	9.28	10.48	9.11	*	+
	NT2RP2006090	6.62	6.62	5.78	3.27	3.55	3.64	**	-
	NT2RP2006092	3.78	3.78	8.3	6.18	8.04	7.07		
10	NT2RP2006097	14.05	14.05	40.38	31.2	25.81	40.02		
	NT2RP2006098	1.94	1.94	4.27	4.52	4.61	7.65		
	NT2RP2006099	3.84	3.84	11.02	10.65	10.99	13.34		
15	NT2RP2006100	2.87	2.87	5.78	3.63	7.31	5.19		
	NT2RP2006103	2.39	2.39	5.54	2.6	3.93	1.71		
	NT2RP2006106	6.48	6.48	21.51	18.05	24.81	22.3		
	NT2RP2006127	3.17	3.17	4.92	1.62	1.26	1.21	*	-
20	NT2RP2006134	4.25	4.25	4.41	6.08	6.7	5.47	**	+
	NT2RP2006141	3.91	3.91	7.94	7.45	6.04	9.08		
	NT2RP2006166	3.1	3.1	10.65	9.01	8.94	7.85		
25	NT2RP2006176	2.15	2.15	4.26	3.95	5.73	4.69		
	NT2RP2006181	1.68	1.68	2.84	3.21	3.14	2.45		
	NT2RP2006184	8.85	8.85	17.16	20.8	19.95	17.1		
	NT2RP2006186	3.01	3.01	4.57	2.77	2.29	4.33		
30	NT2RP2006196	5.24	5.24	7.21	5.25	5.23	4.16		
	NT2RP2006199	5.06	5.06	4.38	3.81	3.65	3.64	**	-
	NT2RP2006200	0.87	0.87	3.43	4.37	4.52	2.17		
35	NT2RP2006210	20.08	20.08	59.85	75.37	70.55	96.59	*	+
	NT2RP2006219	2.88	2.88	6.26	5.97	5.11	7.36		
	NT2RP2006224	3.7	3.7	7.55	9	7.7	8.93		
	NT2RP2006237	1.97	1.97	4.79	3.45	2.74	4.14		
40	NT2RP2006238	3.9	3.9	6.33	4.2	4.69	3.93		
	NT2RP2006258	4.5	4.5	6.73	3.07	4.27	4.39		
	NT2RP2006261	7.32	7.32	3.98	2.04	3.19	7.69		
45	NT2RP2006269	4.11	4.11	7.96	9.52	5.46	9.06		
	NT2RP2006275	3.67	3.67	30.36	23.46	35.36	25.14		
	NT2RP2006282	3.16	3.16	8.89	8.85	8.4	7.05		
	NT2RP2006302	5.69	5.69	12.68	13.12	12.4	11.87		
50	NT2RP2006312	4.88	4.88	8.22	8.47	9.13	9.8		
	NT2RP2006320	4.27	4.27	9.87	6.42	9.32	9.69		
	NT2RP2006321	3.27	3.27	4.23	2.79	4.99	4.13		
55	NT2RP2006323	4.1	4.1	2.59	2.39	3.6	1.83		
	NT2RP2006333	0.67	0.67	1.82	1.7	1.04	1.17		

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	NT2RP2005789	3.98	3.98	11.24	7.89	9.52	8.86		
	NT2RP2005799	2.45	2.45	6.35	2.64	4.67	3.7		
5	NT2RP2005804	9.01	9.01	25	27.85	27.32	30.57		
	NT2RP2005812	2.63	2.63	4.83	2.9	3.89	5.21		
	NT2RP2005815	2.48	2.48	3.15	2.38	3.21	5.2		
10	NT2RP2005835	5.99	5.99	11.26	7.13	13.74	11.69		
	NT2RP2005841	2.32	2.32	10.04	4.89	7.43	11.23		
	NT2RP2005853	1.29	1.29	4.44	2.71	4.6	4.96		
	NT2RP2005857	7.37	7.37	9.87	13.46	7.93	20.27		
15	NT2RP2005859	2.76	2.76	5	2.91	6.14	4.78		
	NT2RP2005860	1.41	1.41	3.54	1.45	1.89	2.22		
	NT2RP2005863	3.03	3.03	6.55	10.76	18.29	15.94	*	+
	NT2RP2005868	3.86	3.86	5.85	5.1	6.3	7.77		
20	NT2RP2005876	5.7	5.7	12.31	7.84	8.29	8.2		
	NT2RP2005878	2.26	2.26	8.44	5.25	4.95	6.32		
	NT2RP2005883	13.54	13.54	21.06	23.75	9.57	28.09		
25	NT2RP2005886	7.18	7.18	50.05	51.13	62.09	50.14		
	NT2RP2005887	3.76	3.76	6.51	4.74	8.05	4.81		
	NT2RP2005890	4.17	4.17	9.77	11.87	17.13	12.15	*	+
	NT2RP2005901	3.19	3.19	5.69	3.91	6.18	5.14		
30	NT2RP2005902	3.17	3.17	4.33	4.78	4.77	6.25		
	NT2RP2005908	3.09	3.09	7.86	4.89	3.7	6.34		
	NT2RP2005927	1.77	1.77	2.66	2.25	4.36	3.07		
35	NT2RP2005933	2.5	2.5	5.59	6.4	4.77	6.42		
	NT2RP2005941	2.09	2.09	5.2	3.31	4.41	3.9		
	NT2RP2005942	4	4	6.86	3.08	4.59	4.64		
	NT2RP2005946	4.63	4.63	9.49	5.33	7.06	6.24		
40	NT2RP2005970	5.44	5.44	14	16.16	22.05	18.9	*	+
	NT2RP2005980	3.71	3.71	5.25	2.69	3.46	2.37		
	NT2RP2005994	2.99	2.99	6.76	4.28	3.28	5.14		
45	NT2RP2006004	1.31	1.31	2.89	2.07	6.09	2.58		
	NT2RP2006013	1.38	1.38	4.91	3.1	5.07	4.92		
	NT2RP2006023	8.37	8.37	17.77	20	21.43	21.14	*	+
	NT2RP2006028	5.03	5.03	10.23	7.47	9.89	9.71		
50	NT2RP2006038	4.67	4.67	5.86	2.79	5.4	1.09		
	NT2RP2006042	8.3	8.3	7.22	6.63	5.89	6.3	*	-
	NT2RP2006043	5.65	5.65	7.59	7.6	10.99	8.29		
55	NT2RP2006052	1.48	1.48	4.48	4.13	3.12	4.54		
	NT2RP2006057	3.73	3.73	6.23	5.69	3.83	4.95		

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	NT2RP2005614	3.06	3.06	5.62	3.68	4.11	2.45		
	NT2RP2005620	3.47	3.47	6.26	3.6	3.92	3.11		
5	NT2RP2005622	6.14	6.14	5.07	6.21	7.43	4.61		
	NT2RP2005632	5.72	5.72	10.95	11.57	10.42	14.89		
	NT2RP2005635	2.22	2.22	19.06	18.14	23.77	18.14		
	NT2RP2005637	1.53	1.53	8	3.73	3.71	4.14		
10	NT2RP2005640	1.72	1.72	7.22	7.49	8.73	6.06		
	NT2RP2005645	4.68	4.68	11.8	10.61	11.47	9.67		
	NT2RP2005651	3.45	3.45	7.88	7.64	6.78	10.15		
15	NT2RP2005654	4.08	4.08	4.14	3.02	2.52	3.8		
	NT2RP2005666	4.91	4.91	5.27	4.34	7.7	4.74		
	NT2RP2005669	7.15	7.15	7.95	7.05	11.14	8.21		
	NT2RP2005670	2.35	2.35	6.91	7.77	5.04	5.2		
20	NT2RP2005671	3.12	3.12	7.83	10.77	8.9	9.78	*	+
	NT2RP2005675	7.32	7.32	37.84	34.46	40.94	40.02		
	NT2RP2005683	2.56	2.56	7.01	7.16	5.19	7.16		
25	NT2RP2005690	2.84	2.84	4.48	2.82	3.74	3.4		
	NT2RP2005694	4.07	4.07	5.49	3.77	6.26	3.54		
	NT2RP2005701	5.97	5.97	8.82	10.39	10.35	9.52	*	+
	NT2RP2005712	5.67	5.67	5.28	4.83	7.94	6.33		
30	NT2RP2005719	1.86	1.86	3.26	4.42	3.8	3.76	*	+
	NT2RP2005722	4.16	4.16	11.13	13.39	15.7	15.94	*	+
	NT2RP2005723	2.71	2.71	4.2	3.65	4.58	3.67		
35	NT2RP2005726	2.55	2.55	4.13	2.86	4.01	3.22		
	NT2RP2005729	4.64	4.64	9.94	10.21	10.7	10.62		
	NT2RP2005731	3.05	3.05	3.39	2.51	2.16	1.27	*	-
	NT2RP2005732	9.41	9.41	57.73	48.37	75.21	41.64		
40	NT2RP2005737	10.75	10.75	22.28	27.16	25.02	17.59		
	NT2RP2005741	3.03	3.03	5.35	3.68	3.31	3.37		
	NT2RP2005748	1.86	1.86	5.94	3.8	3.72	2.95		
45	NT2RP2005752	2.46	2.46	5.55	3.27	4.37	3.8		
	NT2RP2005753	8.45	8.45	14.76	11.12	11.69	14.45		
	NT2RP2005763	3	3	8.03	4.22	4.77	5		
	NT2RP2005767	3.72	3.72	7.79	5.55	6.75	6.29		
50	NT2RP2005773	8.11	8.11	10.02	10.6	11.59	13.37	*	+
	NT2RP2005774	4.25	4.25	12.72	6.86	12.24	14.61		
	NT2RP2005775	3.75	3.75	7.2	3.35	4.83	5.63		
	NT2RP2005781	5.11	5.11	9.88	9.82	7.19	12.94		
55	NT2RP2005784	5.41	5.41	11.51	7.68	12.12	14.06		

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	NT2RP2005419	2.65	2.65	9.05	6.37	8.5	6.77		
	NT2RP2005425	5.63	5.63	18.38	15.27	18.89	15.46		
5	NT2RP2005429	3.23	3.23	-5.85	5.41	6.65	5.64		
	NT2RP2005436	4.65	4.65	10.5	7.02	4.28	4.97		
	NT2RP2005441	2.28	2.28	5.62	3.36	3.77	5.79		
	NT2RP2005442	24.92	24.92	40.66	34.62	25.56	41.66		
10	NT2RP2005444	10.72	10.72	19.24	21.92	21.07	25.56	*	+
	NT2RP2005453	2.79	2.79	7.44	2.63	4.09	3.15		
	NT2RP2005457	15.12	15.12	23.21	28.69	37.38	31.61	*	+
15	NT2RP2005458	2.47	2.47	5.27	3.55	4.16	4.95		
	NT2RP2005463	7.73	7.73	15.23	15.65	22.11	25.05	*	+
	NT2RP2005464	5.96	5.96	11.91	9.22	4.67	10.35		
	NT2RP2005465	1.81	1.81	6.69	3.86	3.75	3.74		
20	NT2RP2005472	10.98	10.98	32.59	28.21	27.9	25.85		
	NT2RP2005476	5.01	5.01	8.99	7.01	6.98	6.08		
	NT2RP2005490	7.51	7.51	21.09	18.18	25.55	23.45		
25	NT2RP2005491	4.99	4.99	12.47	8.63	10.12	8.78		
	NT2RP2005495	3.56	3.56	5.77	3.38	4.55	4.3		
	NT2RP2005496	4.84	4.84	18.25	11.3	13.16	11.28		
	NT2RP2005498	2.92	2.92	7.45	5.18	5.03	4.98		
30	NT2RP2005501	2.04	2.04	5.54	3.12	4.34	2.46		
	NT2RP2005506	124.3	124.3	217.82	139.27	121.83	104.81		
	NT2RP2005509	6.97	6.97	10.45	11.4	9.61	15.73		
35	NT2RP2005514	3.93	3.93	6	4.06	7.05	4.39		
	NT2RP2005520	14.95	14.95	32.39	27.11	39.97	33.03		
	NT2RP2005525	6.19	6.19	7.01	7.81	7.68	4.79		
	NT2RP2005531	2.18	2.18	3.33	1.67	2.12	1.9		
40	NT2RP2005535	4.66	4.66	9.09	9.34	7.79	8.91		
	NT2RP2005539	3.39	3.39	6.22	6.43	5.84	7.45		
	NT2RP2005540	3.2	3.2	7.15	4.79	5.58	6.59		
45	NT2RP2005541	21.25	21.25	39.57	25.85	38.31	39.61		
	NT2RP2005549	2.69	2.69	7.66	6.72	4.85	7.11		
	NT2RP2005555	7.97	7.97	10.1	14.96	16.19	15.37	**	+
	NT2RP2005557	4.89	4.89	8.47	4.03	6.52	6.26		
50	NT2RP2005581	3.93	3.93	9.61	6.32	7.95	6.89		
	NT2RP2005586	1.56	1.56	3.18	3.21	2.92	4.74		
	NT2RP2005597	2.77	2.77	2.93	2.98	4.1	3.84		
55	NT2RP2005600	1.81	1.81	3.71	4.03	4.29	4.44	*	+
	NT2RP2005605	4.93	4.93	14.29	13.17	15.14	15.75		

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	NT2RP2005126	5.88	5.88	8.31	8.9	14.96	8.51		
	NT2RP2005135	5.08	5.08	5.22	4.65	6.59	5.47		
5	NT2RP2005139	1.94	1.94	-2.77	1.87	1.81	2.45		
	NT2RP2005140	3.82	3.82	4.86	12.39	6.72	8.55	*	+
	NT2RP2005144	4.04	4.04	6.31	6.69	5.97	9.7		
	NT2RP2005147	2.23	2.23	5.49	5.61	6.15	6.25		
10	NT2RP2005148	2.86	2.86	5.63	3.83	6.65	4.83		
	NT2RP2005159	3.92	3.92	5.6	4.94	6.38	7.41		
	NT2RP2005162	3.23	3.23	5.56	4.57	5.4	4.21		
15	NT2RP2005163	9.15	9.15	20.61	24.53	28.92	23.77	*	+
	NT2RP2005168	2.87	2.87	6.14	5.24	4.79	4.88		
	NT2RP2005181	2.64	2.64	5.42	3.4	2.11	1.98		
	NT2RP2005204	5.4	5.4	7.81	9.08	11.94	11.81	*	+
20	NT2RP2005219	4.61	4.61	9.64	7.09	10.28	8.7		
	NT2RP2005227	3.59	3.59	10.43	7.55	5.36	9.97		
	NT2RP2005237	26.49	26.49	94.81	86.96	105.8	93.92		
25	NT2RP2005239	2.24	2.24	6.07	2.62	4.27	4.34		
	NT2RP2005247	10.63	10.63	37.59	35.58	46.1	46.95		
	NT2RP2005254	4.35	4.35	9.14	5.7	6.44	6.93		
	NT2RP2005270	9.06	9.06	17.44	10.82	9.28	17.11		
30	NT2RP2005276	7.19	7.19	11.53	10.88	11.68	15.71		
	NT2RP2005287	7.98	7.98	11.97	8.37	7.7	13.36		
	NT2RP2005288	2.51	2.51	5.14	2.89	5.59	5.22		
35	NT2RP2005289	4.26	4.26	8.48	6.68	9.08	7.49		
	NT2RP2005293	5	5	6.93	13.68	14.37	15.66	**	+
	NT2RP2005315	5.79	5.79	10.64	8.04	12.95	16.68		
	NT2RP2005322	5.05	5.05	15.42	18.91	11.33	22.43		
40	NT2RP2005325	8.45	8.45	18.4	15.57	13.63	20.01		
	NT2RP2005336	1.71	1.71	6.68	4.18	5.74	5.3		
	NT2RP2005343	2.44	2.44	7.48	3.91	4.11	5.89		
45	NT2RP2005344	3.39	3.39	4.83	2.37	2.67	3.32		
	NT2RP2005347	3.14	3.14	3.61	3.34	2.96	3.53		
	NT2RP2005354	6.49	6.49	11.79	10.37	13	11.38		
	NT2RP2005358	35.87	35.87	109.04	101.37	134.72	117.96		
50	NT2RP2005360	2.93	2.93	5	3.59	4.97	3.84		
	NT2RP2005378	5.27	5.27	13.12	7.54	8.7	13.93		
	NT2RP2005391	3.06	3.06	5.41	4.21	6.76	7.72		
55	NT2RP2005393	1.61	1.61	6.34	4.86	6.16	4.07		
	NT2RP2005407	2.59	2.59	5.71	4.28	5.65	4.64		

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	NT2RP2004810	3.44	3.44	8.83	7.37	7.84	6.03		
	NT2RP2004816	5.58	5.58	12.1	11.22	8.76	11.15		
5	NT2RP2004837	4.13	4.13	9.89	10.43	7.23	12.98		
	NT2RP2004841	0.91	0.91	2.86	3.69	4.03	8.87		
	NT2RP2004847	3.25	3.25	13.75	13.82	13.87	17.16		
	NT2RP2004861	2.3	2.3	5.23	2.33	4.23	2.46		
10	NT2RP2004897	3.35	3.35	6.43	4.26	3.27	3.35		
	NT2RP2004932	6.64	6.64	10.16	7.96	8.53	6.91		
	NT2RP2004933	4.63	4.63	3.41	2.98	2.93	3.2	*	-
15	NT2RP2004936	3.69	3.69	6.41	4.56	4.42	7.53		
	NT2RP2004951	2.98	2.98	10.48	5.09	5.22	19.28		
	NT2RP2004959	3.13	3.13	6.61	6.43	6.26	6.5		
	NT2RP2004961	2.1	2.1	4.79	4.89	6.49	5.44		
20	NT2RP2004962	2.27	2.27	7.28	4.5	5.57	4.47		
	NT2RP2004966	2.26	2.26	6.07	4.1	4.1	2.97		
	NT2RP2004967	3.87	3.87	6.16	4.07	4.82	3.26		
25	NT2RP2004974	5.27	5.27	5.43	3.59	3.47	3.89	**	-
	NT2RP2004978	2.68	2.68	5.26	4.17	6.39	5.09		
	NT2RP2004982	0.57	0.57	1.82	2.2	1.94	1.72		
	NT2RP2004985	16.03	16.03	45.34	44.65	46.12	54.4		
30	NT2RP2004999	2.21	2.21	5.64	4.27	8.86	10.34		
	NT2RP2005000	3.62	3.62	5.76	4.33	4.76	4.65		
	NT2RP2005001	5.41	5.41	7.91	8.26	9.15	8.32		
35	NT2RP2005003	3.8	3.8	7.2	6.11	7.91	6.2		
	NT2RP2005012	6.61	6.61	20.14	18.41	20.96	17.87		
	NT2RP2005018	1.9	1.9	4.24	3.29	2.24	2.91		
	NT2RP2005020	6.12	6.12	23.58	19.97	19.94	22.96		
40	NT2RP2005022	1.65	1.65	5.01	5.09	7.24	4.77		
	NT2RP2005027	5.96	5.96	38.61	42.51	40.08	33.77		
	NT2RP2005031	1.54	1.54	4.99	3.94	4.53	3.92		
45	NT2RP2005035	44.19	44.19	94.82	116.52	107.36	106.69	*	+
	NT2RP2005037	4.28	4.28	5.87	7.91	10.26	7.09	*	+
	NT2RP2005038	4.86	4.86	4.84	1.85	2.29	3.1	**	-
	NT2RP2005048	9.92	9.92	30.91	30.14	33.38	32.48		
50	NT2RP2005069	16.01	16.01	34.88	21.99	20.63	27.64		
	NT2RP2005073	7.36	7.36	30.35	29.36	28.24	30.33		
	NT2RP2005097	2.39	2.39	5.6	5.27	5.2	4.6		
55	NT2RP2005108	1.76	1.76	3.95	2.84	4.21	7.12		
	NT2RP2005116	3.53	3.53	5.96	6.27	5.42	5.89		

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	NT2RP2004512	4.25	4.25	7.62	5.84	6.41	7.12		
	NT2RP2004523	5.18	5.18	11.04	8.44	7.56	10.21		
5	NT2RP2004524	2.19	2.19	-6.33	4.97	5.32	5.56		
	NT2RP2004536	8.99	8.99	16.96	14.1	14.51	17.51		
	NT2RP2004538	8.03	8.03	24.44	20.15	25.59	22.24		
10	NT2RP2004548	4.45	4.45	9.92	7.39	9.1	10.51		
	NT2RP2004551	4.95	4.95	5.62	7.17	9.98	6.21		
	NT2RP2004556	83.73	83.73	210.17	226.48	298.92	241.84		
	NT2RP2004568	5.19	5.19	11.18	6.52	9.16	9.99		
15	NT2RP2004580	3.98	3.98	7.71	5.71	7.88	7.04		
	NT2RP2004585	11.28	11.28	49.82	36.69	46.91	64.56		
	NT2RP2004587	1.85	1.85	4.16	2.07	2.89	3		
20	NT2RP2004594	4.56	4.56	9.24	11.47	12.21	28.18		
	NT2RP2004600	3.49	3.49	5.76	2.22	3.22	3.09		
	NT2RP2004602	4.62	4.62	6.32	6	8.49	6.26		
	NT2RP2004606	392.21	392.21	581.19	612.4	897.5	764.63	*	+
25	NT2RP2004614	2.92	2.92	4.73	2.69	3.63	3.81		
	NT2RP2004648	2.52	2.52	4.96	3.24	5.01	4.12		
	NT2RP2004655	5.69	5.69	10.1	8.37	6.76	9.46		
30	NT2RP2004664	3.64	3.64	5.35	3.59	4.62	5.97		
	NT2RP2004670	1.98	1.98	3.81	1.98	3.71	4.27		
	NT2RP2004675	3.37	3.37	9.29	4.08	5.87	5.33		
	NT2RP2004681	3.46	3.46	7.56	5.72	8.92	7.55		
35	NT2RP2004689	2.63	2.63	5.75	5.75	4.73	7.87		
	NT2RP2004709	3.93	3.93	7.79	4.46	2.89	5.25		
	NT2RP2004710	3.15	3.15	8.37	5.63	4.61	6.88		
40	NT2RP2004721	1.79	1.79	5.99	3.39	4.41	2.78		
	NT2RP2004736	3.26	3.26	5.81	6.11	4.79	4.63		
	NT2RP2004743	4.94	4.94	7.96	5.94	6.67	7.36		
	NT2RP2004750	6.21	6.21	17.46	11.9	15.49	11.01		
45	NT2RP2004755	11.65	11.65	19.9	14.84	22.87	19.91		
	NT2RP2004767	3.54	3.54	9	4.05	5.8	4.81		
	NT2RP2004768	3.48	3.48	29.51	18.48	18.73	19.6		
	NT2RP2004775	4.68	4.68	5.68	7.71	5.62	8.26		
50	NT2RP2004791	7.23	7.23	16.58	9.33	10.24	11.68		
	NT2RP2004794	14.01	14.01	25.74	23.04	16.86	22.78		
	NT2RP2004795	5.15	5.15	7.97	6.96	5.67	11.2		
55	NT2RP2004799	6.74	6.74	10.99	5.35	8.58	6.3		
	NT2RP2004802	6.35	6.35	11.79	6.1	7.62	6.24		

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	NT2RP2004196	4.28	4.28	13.77	10.02	7.87	14.61		
	NT2RP2004205	2.67	2.67	8.14	8.64	6.62	7.81		
5	NT2RP2004207	2.57	2.57	-4.38	4.15	4.97	3.59		
	NT2RP2004226	2.09	2.09	4.95	4.11	6.15	5.33		
	NT2RP2004232	2.79	2.79	6.52	6	6.59	5.33		
10	NT2RP2004239	3.57	3.57	4.49	2.71	3.97	5.6		
	NT2RP2004240	7.07	7.07	12.57	13	15.8	8.95		
	NT2RP2004242	3.87	3.87	6.52	5.77	6.94	7.27		
	NT2RP2004245	1.74	1.74	3.47	2.42	3.29	3.15		
15	NT2RP2004270	9.77	9.77	33.78	28.39	27.43	29.48		
	NT2RP2004300	2	2	5.22	4.34	4.52	3.26		
	NT2RP2004304	6.46	6.46	15.37	17.41	12.33	13.9		
20	NT2RP2004313	3.17	3.17	3.78	5.51	4.18	4.63	*	+
	NT2RP2004316	3.46	3.46	5.84	4.9	4.96	4.04		
	NT2RP2004321	4.71	4.71	6.06	6.79	7.43	6.29	*	+
	NT2RP2004336	4.19	4.19	4.97	2.73	4.28	4.53		
25	NT2RP2004339	5.3	5.3	20.89	17.11	18.07	15.39		
	NT2RP2004347	1.39	1.39	3.99	4.78	5	4.19		
	NT2RP2004364	2.26	2.26	6.52	5.08	6.72	4.76		
30	NT2RP2004365	3.18	3.18	6.58	6.68	6.34	7.7		
	NT2RP2004366	2.49	2.49	6.06	4.49	4.71	3.08		
	NT2RP2004373	8.17	8.17	14.38	7.1	7.22	5.91		
	NT2RP2004375	9.27	9.27	13.98	20.89	26.85	20.68	**	+
35	NT2RP2004389	5.25	5.25	5.62	5.01	6.26	5.61		
	NT2RP2004392	8.88	8.88	23.7	13.04	19.48	20.89		
	NT2RP2004396	1.98	1.98	6.27	6.65	4.98	6.17		
	NT2RP2004399	5.24	5.24	8.12	12.56	7.74	9.52		
40	NT2RP2004400	2.07	2.07	3.55	2.36	3.47	2		
	NT2RP2004404	15.79	15.79	46	45.56	40.89	41.74		
	NT2RP2004410	16.64	16.64	24.04	27.99	33.46	32.69	*	+
45	NT2RP2004412	5.84	5.84	6.74	7.37	9.71	7.37		
	NT2RP2004414	4.27	4.27	5.09	3.81	4.89	3.8		
	NT2RP2004425	3.71	3.71	6.53	3.73	3.18	4.04		
	NT2RP2004447	1.93	1.93	5.68	2.75	5.22	3.56		
50	NT2RP2004463	13.57	13.57	16.23	16.84	16.25	20.26		
	NT2RP2004476	9.11	9.11	12.69	11.89	12.66	15.87		
	NT2RP2004488	3.82	3.82	8.52	4.59	6.02	5.37		
55	NT2RP2004490	2.88	2.88	3.86	2.31	2.96	4.1		
	NT2RP2004495	35.59	35.59	88.76	96.31	109.31	123.5	*	+

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	NT2RP2003912	3.83	3.83	14.48	5.91	7.43	6.74		
	NT2RP2003931	1.81	1.81	6.03	3.95	6.86	4.42		
5	NT2RP2003940	2.31	2.31	-9.51	7.1	6.2	7.3		
	NT2RP2003950	2.81	2.81	5.48	3.84	5.57	2.98		
	NT2RP2003952	1.86	1.86	5.63	2.58	4.23	2.98		
	NT2RP2003968	4.82	4.82	7.38	9.86	11.76	13.51	*	+
10	NT2RP2003976	5.35	5.35	9.56	12.56	12.6	13.2	*	+
	NT2RP2003981	3.27	3.27	7.41	4.62	2.03	4.07		
	NT2RP2003984	5.57	5.57	15.87	10.21	4.25	10.34		
15	NT2RP2003986	2.79	2.79	6.22	6.29	6.32	5.17		
	NT2RP2003988	2.36	2.36	6.84	4.51	7.42	5		
	NT2RP2004013	8.46	8.46	13.75	14.68	13.19	17		
	NT2RP2004014	4.24	4.24	10.07	4.06	5.12	4.08		
20	NT2RP2004036	6.88	6.88	14.85	14.08	19.02	16.03		
	NT2RP2004041	2.77	2.77	5.02	3.96	4.43	5.19		
	NT2RP2004042	1.99	1.99	4.6	4.41	2.02	4.09		
25	NT2RP2004049	4.68	4.68	19.13	14.24	15.5	16.3		
	NT2RP2004060	5.7	5.7	10.41	7.09	8.67	10.84		
	NT2RP2004066	2.17	2.17	4.31	3.05	4.83	3.65		
	NT2RP2004069	3.99	3.99	7.24	3.54	6	4.26		
30	NT2RP2004076	3.73	3.73	5.82	1.92	4.2	3.61		
	NT2RP2004080	4.21	4.21	9.26	4.45	6.47	6.15		
	NT2RP2004081	3.27	3.27	5.39	3.51	3.71	4.5		
35	NT2RP2004098	2.32	2.32	6.48	5.4	3.1	5.75		
	NT2RP2004108	3.82	3.82	9.56	7.18	5.89	7.56		
	NT2RP2004124	3.13	3.13	5.9	3.68	5.82	3.92		
	NT2RP2004130	3.67	3.67	9.32	5.51	9.12	8.4		
40	NT2RP2004133	2.05	2.05	6.41	3.69	6.54	6.25		
	NT2RP2004141	5.72	5.72	7.15	5.14	7.05	7.05		
	NT2RP2004142	5.33	5.33	8.1	4.18	5.45	3.93		
45	NT2RP2004152	3.34	3.34	4.78	5.7	7.49	4.39		
	NT2RP2004165	3.71	3.71	8.3	5.87	5.92	6.54		
	NT2RP2004170	1.86	1.86	5.97	5.37	4.17	4.94		
	NT2RP2004172	2.93	2.93	5.24	4.69	5.58	4.26		
50	NT2RP2004176	3.45	3.45	8.4	7.77	10.21	8.98		
	NT2RP2004179	4.01	4.01	9.17	3.94	5.07	4.15		
	NT2RP2004187	3.16	3.16	6.36	3.87	3.88	4.59		
55	NT2RP2004190	5.1	5.1	5.46	5.49	7.33	9.98		
	NT2RP2004194	7.54	7.54	14.57	18.5	23.44	19.83	*	+

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	NT2RP2003596	4.86	4.86	10.86	14.43	13.12	17.96	*	+
	NT2RP2003599	6.49	6.49	12.46	14.29	10.17	11.98		
5	NT2RP2003600	1.88	1.88	2.95	3.02	3.64	6.36		
	NT2RP2003604	7.09	7.09	8.97	16.39	13.03	16.68	**	+
	NT2RP2003629	3.72	3.72	5.25	3.11	4.56	2.38		
	NT2RP2003630	4.09	4.09	6.66	4.79	6.78	3.84		
10	NT2RP2003643	5.49	5.49	4.88	7.15	9.8	8.51	*	+
	NT2RP2003655	4.27	4.27	11.12	7.52	6.38	7.59		
	NT2RP2003664	12.29	12.29	24.31	17.9	18.07	17.11		
15	NT2RP2003668	2.52	2.52	5.01	3.3	3.18	3.62		
	NT2RP2003687	1.61	1.61	2.77	1.63	2.42	1.71		
	NT2RP2003691	3.03	3.03	5.07	3.7	4.21	4.57		
	NT2RP2003702	3.99	3.99	6.14	2.89	3.02	2.89		
20	NT2RP2003704	3.31	3.31	4.12	2.65	3.84	1.99		
	NT2RP2003706	2.44	2.44	1.24	1.72	1.42	1.6		
	NT2RP2003713	4.11	4.11	5.49	4.16	3.89	3.87		
25	NT2RP2003714	3.39	3.39	7.8	5.19	5.31	5.32		
	NT2RP2003727	3.96	3.96	11.63	4.81	6.08	10.46		
	NT2RP2003737	2.52	2.52	8.58	4.88	6.47	4.6		
	NT2RP2003751	1.66	1.66	4.84	1.24	1.67	1.37		
30	NT2RP2003760	2.52	2.52	5.47	3.87	4.19	5.45		
	NT2RP2003764	2.1	2.1	2.81	1.44	1.87	4.62		
	NT2RP2003769	5.52	5.52	11.45	7.41	8.17	10.64		
35	NT2RP2003770	7.43	7.43	12.42	7.17	5.67	12.82		
	NT2RP2003777	3.44	3.44	6.78	5.01	5.6	9.57		
	NT2RP2003781	4.93	4.93	15.85	13.04	11.91	13.48		
	NT2RP2003785	9.69	9.69	13.44	11.1	10.68	8.99		
40	NT2RP2003793	9.32	9.32	9.5	8.29	13.22	10.51		
	NT2RP2003806	5.6	5.6	12.03	8.54	8.75	12.97		
	NT2RP2003825	10.73	10.73	62.01	57.88	71.84	82.78		
45	NT2RP2003840	3.19	3.19	6.07	3.86	4.44	4.71		
	NT2RP2003857	4.02	4.02	4.94	3.15	4.61	6.12		
	NT2RP2003859	1.82	1.82	6	3.35	4.16	4.21		
	NT2RP2003871	5.22	5.22	9.43	5.1	4.59	7.79		
50	NT2RP2003876	3.82	3.82	8.8	5.92	5.87	7.92		
	NT2RP2003878	3.38	3.38	6.49	3.8	4.9	4.11		
	NT2RP2003885	2.46	2.46	3.09	1.66	3.29	2.37		
55	NT2RP2003898	5.39	5.39	8.91	12.3	12.73	18.25	*	+
	NT2RP2003902	5.09	5.09	10.78	8.24	8.23	10.42		

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	NT2RP2003390	9.1	9.1	17.93	16.66	14.27	12.94		
	NT2RP2003391	9.39	9.39	12.9	11.96	9.91	12.84		
5	NT2RP2003393	4.23	4.23	-6.99	6.14	5.03	9.44		
	NT2RP2003394	8.67	8.67	16.21	17.56	21.75	17.23		
	NT2RP2003401	4.39	4.39	5.97	3.52	3.72	2.9		
10	NT2RP2003403	3.42	3.42	7.64	6.62	8.55	7.79		
	NT2RP2003433	3.02	3.02	15.54	13.62	13.5	13.08		
	NT2RP2003445	3.2	3.2	4.74	4.08	3.5	3.93		
	NT2RP2003446	2.67	2.67	6.23	5.06	6.05	4.47		
15	NT2RP2003456	2.04	2.04	6.57	4.26	5.89	3.81		
	NT2RP2003466	3.56	3.56	20.09	17.34	25.96	23.53		
	NT2RP2003469	6.2	6.2	5.65	6.19	7.53	6.33		
20	NT2RP2003470	5.64	5.64	6.47	5.06	6.11	6.44		
	NT2RP2003471	2.72	2.72	3.88	3.22	2.78	3.38		
	NT2RP2003480	7.15	7.15	20.74	19.77	19.89	21.14		
	NT2RP2003495	3.99	3.99	6.03	8.07	7.24	10.72	*	+
25	NT2RP2003499	1.52	1.52	4.58	3.67	4.05	2.75		
	NT2RP2003505	0.98	0.98	3.21	2.62	3.88	1.4		
	NT2RP2003506	2.54	2.54	6.53	5.65	5.36	4.78		
30	NT2RP2003511	3.67	3.67	5.57	4.22	3.1	2.96		
	NT2RP2003513	3.79	3.79	6.01	5.49	5.57	5.71		
	NT2RP2003517	2.9	2.9	2.52	1.32	1.11	0.85	**	-
	NT2RP2003522	11.08	11.08	19.77	10.55	11.42	16.52		
35	NT2RP2003525	5.12	5.12	14.93	12.19	10.72	11.79		
	NT2RP2003533	3.36	3.36	10.44	12.12	10.72	12.94		
	NT2RP2003541	6.72	6.72	11.29	12.02	13.42	11.7		
40	NT2RP2003543	2.48	2.48	5.96	4.17	3.55	6.54		
	NT2RP2003545	2.59	2.59	4.85	2.22	3.6	1.85		
	NT2RP2003559	4.92	4.92	4.81	3.97	3.84	3.37	**	-
	NT2RP2003564	4.46	4.46	3.93	2.53	1.97	2.42	**	-
45	NT2RP2003565	4.94	4.94	50.48	41.12	48.32	37.82		
	NT2RP2003567	3.51	3.51	16.65	16.25	19.43	16.05		
	NT2RP2003575	4.44	4.44	18.78	19.56	22.63	20.7		
	NT2RP2003576	102.12	102.12	203.44	206.62	128.42	171.89		
50	NT2RP2003579	11.45	11.45	26.58	38.62	39.51	39.88	*	+
	NT2RP2003581	3.85	3.85	6.1	4.33	4.38	3.96		
	NT2RP2003587	8.37	8.37	11.47	13.35	14.11	12.14	*	+
55	NT2RP2003590	7.15	7.15	9.08	11.06	13.15	14.91	*	+
	NT2RP2003593	1.58	1.58	4.57	7.84	4.43	8.59		

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	NT2RP2003150	1.45	1.45	5.71	3.98	5.2	4.3		
	NT2RP2003157	6.93	6.93	34.27	34.29	31.85	32.84		
5	NT2RP2003158	6.3	6.3	-25.32	26.87	28.69	59.31		
	NT2RP2003161	2.73	2.73	3.36	2.51	2.82	6.12		
	NT2RP2003164	1.96	1.96	2.1	1.28	1.87	2.46		
10	NT2RP2003165	2.18	2.18	5.94	3.1	3.69	4.84		
	NT2RP2003177	1.63	1.63	4.37	2.79	3.03	4.42		
	NT2RP2003179	1.23	1.23	4.98	4.08	3.63	7.96		
	NT2RP2003194	4.04	4.04	7.2	5.73	6.29	14.77		
15	NT2RP2003206	1.59	1.59	4.47	1.64	3.52	1.44		
	NT2RP2003210	5.06	5.06	15.15	16.14	12.93	15.9		
	NT2RP2003227	1.62	1.62	3.97	2.04	3.66	6.28		
20	NT2RP2003228	6.57	6.57	29.53	29.56	43.94	44.24		
	NT2RP2003230	3.51	3.51	7.91	4.49	8.04	8.46		
	NT2RP2003231	2.22	2.22	5.59	2.46	3.23	3.83		
	NT2RP2003237	2.52	2.52	4.59	4.59	6.4	6.46	*	+
25	NT2RP2003239	2.3	2.3	4.46	2.97	4.46	4.05		
	NT2RP2003243	2.16	2.16	4.13	2.38	3.28	3.98		
	NT2RP2003265	3.93	3.93	5.33	4.22	4.88	4.92		
	NT2RP2003267	2.73	2.73	3.15	3.24	4.17	7.42		
30	NT2RP2003272	6.03	6.03	14.8	16.93	23.85	32.58	*	+
	NT2RP2003277	3.85	3.85	11.29	5.53	8.39	6.39		
	NT2RP2003280	3.47	3.47	9.38	7.67	7.25	6.09		
35	NT2RP2003286	2.18	2.18	4.23	4.13	5	9.61		
	NT2RP2003293	2.98	2.98	6.9	5.66	7.05	7.94		
	NT2RP2003295	4.67	4.67	8.45	8.73	12.39	6.35		
	NT2RP2003297	3.43	3.43	7.57	4.32	6.89	5.28		
40	NT2RP2003300	20.38	20.38	32.04	45.7	53.51	48.07	**	+
	NT2RP2003302	2.88	2.88	4.52	3.46	3.81	7.26		
	NT2RP2003307	0.57	0.57	2.62	1.43	1.49	1.38		
45	NT2RP2003308	1.44	1.44	4.5	2.77	4.44	5.44		
	NT2RP2003311	4.18	4.18	5.83	7.35	4.25	8.2		
	NT2RP2003329	2.99	2.99	4.41	2.63	3.89	4.44		
	NT2RP2003339	3.06	3.06	7.01	3.76	4.92	3.64		
50	NT2RP2003345	4.15	4.15	8.38	2.77	3.97	6.33		
	NT2RP2003347	2.55	2.55	4.23	2.08	1.98	3.46		
	NT2RP2003367	2.15	2.15	4.65	2.7	1.98	1.44		
55	NT2RP2003369	1.34	1.34	4.71	2.16	2.36	0.89		
	NT2RP2003383	4.05	4.05	6.75	7.66	7.17	6.99		

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	NT2RP2002907	2.95	2.95	6.91	5.95	5.5	4.9		
	NT2RP2002925	5.67	5.67	6.73	10.04	11.6	8.45	*	+
5	NT2RP2002927	10.08	10.08	10.63	19.05	21.85	15.48	**	+
	NT2RP2002928	4.32	4.32	4.56	2.65	2.34	2.51	**	-
	NT2RP2002929	3.96	3.96	9.74	8.09	7.86	9.87		
10	NT2RP2002934	1.5	1.5	1.4	2.81	3.01	2.49	**	+
	NT2RP2002939	2.96	2.96	5.09	5.15	6.71	4.91		
	NT2RP2002942	2.4	2.4	5.06	4.35	5.07	10.81		
	NT2RP2002954	5.41	5.41	11.46	7.21	9.1	8.65		
15	NT2RP2002959	8.15	8.15	12.55	13.81	16.28	16.69	*	+
	NT2RP2002974	5.03	5.03	6.53	4.7	3.45	4.54		
	NT2RP2002976	6.92	6.92	17.08	11.84	14.66	12.42		
20	NT2RP2002979	4.41	4.41	8.12	7.03	8.66	7.6		
	NT2RP2002980	6.44	6.44	15.09	15.56	11	17.45		
	NT2RP2002986	3.87	3.87	7.6	6.68	7.4	7.39		
	NT2RP2002987	3.52	3.52	8.23	11.1	9.18	9.4	*	+
25	NT2RP2002988	14.96	14.96	22.92	30.07	31.87	31.36	**	+
	NT2RP2002993	2.97	2.97	4.18	3.8	3.84	2.84		
	NT2RP2003000	4.88	4.88	8.34	6.97	9.62	9.97		
30	NT2RP2003008	4.85	4.85	5.06	3.34	4.76	4.78		
	NT2RP2003020	4.45	4.45	44.26	28.35	46.52	34.33		
	NT2RP2003032	1.91	1.91	4.02	5.82	6.48	6.59	**	+
	NT2RP2003034	4.21	4.21	13.47	13.16	11.15	16.31		
35	NT2RP2003042	2.15	2.15	3.81	4.57	3.65	4.92		
	NT2RP2003050	2.32	2.32	3.56	2.55	2.17	1.83		
	NT2RP2003060	7.27	7.27	15.51	21.53	18.91	17.46	*	+
40	NT2RP2003073	5.61	5.61	8.73	7.06	10.51	8.17		
	NT2RP2003099	5.05	5.05	3.67	3.21	3.73	2.84		
	NT2RP2003108	3.6	3.6	4.23	5.29	3.91	6.62		
	NT2RP2003115	1.68	1.68	5	7.75	4.69	4.84		
45	NT2RP2003117	2.71	2.71	5.69	3.6	4.66	4.13		
	NT2RP2003121	1.83	1.83	3.47	4.03	2.69	3.33		
	NT2RP2003125	4.13	4.13	11.44	15.42	12.55	13.66	*	+
	NT2RP2003127	2.36	2.36	3.94	1.53	1.66	1.75		
50	NT2RP2003129	3.43	3.43	7.09	6.08	6.05	5.42		
	NT2RP2003137	4.49	4.49	6.14	7.58	8.4	6.46	*	+
	NT2RP2003138	4.66	4.66	20.24	16.55	17.45	16.92		
55	NT2RP2003146	6.2	6.2	24.78	18.5	23.25	25.96		
	NT2RP2003148	3.09	3.09	6.73	3.06	4.6	4.04		

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	NT2RP2002602	4.19	4.19	7.53	5.68	8.96	10.42		
	NT2RP2002606	1.27	1.27	2.93	2.26	2.97	3.95		
5	NT2RP2002609	6.12	6.12	9.95	4.79	5.48	7.74		
	NT2RP2002618	2.74	2.74	6.83	4.2	6.34	5.44		
	NT2RP2002621	4.24	4.24	10.22	6.58	7.52	9.79		
	NT2RP2002643	1.79	1.79	4.84	3.11	5.98	3.94		
10	NT2RP2002672	4.48	4.48	9.23	8.03	9.37	9.87		
	NT2RP2002673	4.13	4.13	5.01	8	12.88	17.73	*	+
	NT2RP2002674	2.4	2.4	4.06	2.78	2.37	1.84		
15	NT2RP2002686	2.73	2.73	4.61	3.17	5.19	6.49		
	NT2RP2002688	10.73	10.73	28.07	22.34	33.34	28.71		
	NT2RP2002695	2.62	2.62	7.03	5.26	4.34	5.52		
	NT2RP2002701	7.29	7.29	13.37	11.85	5.18	10.04		
20	NT2RP2002706	3.02	3.02	5.58	6.47	8.14	6.19	*	+
	NT2RP2002710	11.2	11.2	36.97	39.43	33.9	42.75		
	NT2RP2002721	5.53	5.53	9.42	7.33	7.34	8.45		
25	NT2RP2002727	3.56	3.56	6.87	2.17	3.96	3.52		
	NT2RP2002734	3.59	3.59	6.65	5.71	7.65	7.54		
	NT2RP2002736	5.25	5.25	13.13	13.17	13.59	17.58		
	NT2RP2002740	2	2	5.11	3.18	2.81	3.13		
30	NT2RP2002741	2.12	2.12	4.8	4.68	5.39	5.78		
	NT2RP2002750	2.5	2.5	9.22	7.73	9.57	8.18		
	NT2RP2002752	4.59	4.59	10.39	7.93	7.66	8.03		
35	NT2RP2002753	4.49	4.49	9.91	8.66	9.04	14.67		
	NT2RP2002760	4.79	4.79	10.31	3.56	4.5	4.13		
	NT2RP2002769	4.42	4.42	6.06	3.89	5.67	6.43		
	NT2RP2002778	4.13	4.13	10.59	6.8	8.2	9.12		
40	NT2RP2002791	8.89	8.89	54.27	48.75	53.08	50.19		
	NT2RP2002800	1.66	1.66	4.52	4	5.19	5.38		
	NT2RP2002805	3.38	3.38	5.46	4.75	6.44	3.81		
45	NT2RP2002811	3.27	3.27	8.23	5.87	8.13	10.61		
	NT2RP2002824	18.29	18.29	25.05	29.95	34.29	25.05		
	NT2RP2002839	13.26	13.26	31.21	16.09	23.42	16.27		
	NT2RP2002845	5.87	5.87	7.93	4.61	6.12	5.5		
50	NT2RP2002857	2.95	2.95	3.6	2.35	4.23	2.99		
	NT2RP2002862	4.56	4.56	12.49	12.55	9.84	11.34		
	NT2RP2002880	5.27	5.27	13.89	13.5	11.56	11.1		
55	NT2RP2002885	8.6	8.6	17.12	7.56	10.07	10.02		
	NT2RP2002891	1.9	1.9	7.78	4.65	6.63	7.56		

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	NT2RP2002292	5.42	5.42	8.4	6.64	8.08	6.95		
	NT2RP2002299	9.6	9.6	9.22	11.75	19.42	15.97	*	+
5	NT2RP2002304	1.37	1.37	4.78	6.99	5.08	5.52		
	NT2RP2002312	1.21	1.21	2.33	3.78	5.3	3.28	*	+
	NT2RP2002316	3.28	3.28	5.43	7.57	7.21	8.2	**	+
10	NT2RP2002325	1.95	1.95	3.46	2.79	2.22	4.95		
	NT2RP2002333	2.13	2.13	3.03	3.53	4.86	5.69	*	+
	NT2RP2002371	5.43	5.43	9.14	9.72	12.07	11.65	*	+
	NT2RP2002373	10.65	10.65	40.1	36.72	58.84	33.58		
15	NT2RP2002381	4.68	4.68	2.35	2.66	3.19	3.71		
	NT2RP2002385	5.71	5.71	11.84	9.95	11.34	9.47		
	NT2RP2002394	0.94	0.94	1.52	1.24	0.96	1.26		
20	NT2RP2002408	2.7	2.7	5.08	3.89	3.12	4.29		
	NT2RP2002409	3.73	3.73	10.81	10.78	7.95	8.35		
	NT2RP2002424	2.98	2.98	4.22	5.84	6.22	7.85	*	+
	NT2RP2002426	6.44	6.44	11.38	7.59	8.46	8.93		
25	NT2RP2002429	17.2	17.2	24.73	27.87	33.96	20.83		
	NT2RP2002437	4.61	4.61	5.98	4.83	6.47	4.79		
	NT2RP2002439	3.83	3.83	6.69	2.68	3.22	4		
30	NT2RP2002442	13.63	13.63	71.65	57.78	63.05	78.84		
	NT2RP2002457	3.27	3.27	5.31	4.35	4.87	5.82		
	NT2RP2002464	2.17	2.17	5.34	3.29	4.59	4.24		
	NT2RP2002475	3.11	3.11	7.88	5.3	2.83	5.43		
35	NT2RP2002479	3.09	3.09	4.25	1.95	2.99	1.93		
	NT2RP2002487	1.73	1.73	5.15	1.98	2.1	3.04		
	NT2RP2002498	1.52	1.52	2.2	2.62	2.82	4.47		
40	NT2RP2002503	7.63	7.63	31.85	29.32	32.02	31.84		
	NT2RP2002504	3.81	3.81	5.73	6	7.23	11.28		
	NT2RP2002510	2.65	2.65	8.92	4.68	6.59	6.85		
	NT2RP2002520	3.57	3.57	7.17	6.26	8.86	6.61		
45	NT2RP2002527	5.18	5.18	6.02	9	12.37	11.22	**	+
	NT2RP2002533	3.34	3.34	6.27	4.83	6.94	5.88		
	NT2RP2002537	3.22	3.22	4.02	4.09	5.91	10.08		
	NT2RP2002542	4.81	4.81	4.64	5.99	5.93	9.73		
50	NT2RP2002546	4.31	4.31	5.85	6.5	4.91	5.24		
	NT2RP2002549	4.06	4.06	9.33	7.68	10.49	11.65		
	NT2RP2002564	4.11	4.11	11.18	10.67	9.21	9.29		
55	NT2RP2002591	2.45	2.45	7.03	3.31	4.79	5.79		
	NT2RP2002595	9.67	9.67	12.41	12.06	13.39	14.79		

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	NT2RP2002047	2.71	2.71	3.86	2.06	3.87	3.83		
	NT2RP2002050	7.67	7.67	11.66	9.78	6.47	7.77		
5	NT2RP2002052	3.77	3.77	-8.39	6.6	3.99	6.28		
	NT2RP2002058	2.07	2.07	4.11	3.1	4.82	3.39		
	NT2RP2002060	1.48	1.48	3.44	2.1	5.9	3.18		
10	NT2RP2002063	3.61	3.61	5.83	3.3	5.2	6.65		
	NT2RP2002066	10.11	10.11	13.47	5.47	9.88	9.73		
	NT2RP2002070	3.74	3.74	6.3	2.15	3.1	3.83		
	NT2RP2002076	3.72	3.72	5.1	3.35	5.58	3.63		
15	NT2RP2002078	13.09	13.09	105.74	73.87	88.51	76.08		
	NT2RP2002079	5.76	5.76	36.34	31.87	32.66	36.54		
	NT2RP2002099	4.19	4.19	5.82	4.7	6.75	6.03		
20	NT2RP2002105	2.66	2.66	12.25	11.41	14.04	11.23		
	NT2RP2002115	1.63	1.63	5.21	1.98	3.42	1.52		
	NT2RP2002124	3.66	3.66	6	4.56	5.38	3.88		
	NT2RP2002137	3.99	3.99	4.83	2.21	2.1	1.76	**	-
25	NT2RP2002139	24.08	24.08	45.74	51.88	77.99	62.3	*	+
	NT2RP2002154	1.37	1.37	4.13	3.56	2.36	3.2		
	NT2RP2002155	351.63	351.63	869.83	623.53	501.61	620.68		
30	NT2RP2002172	1.5	1.5	3.33	2.78	3.53	5.13		
	NT2RP2002185	3.29	3.29	7.65	7.3	8.56	8.12		
	NT2RP2002188	1.74	1.74	5.95	4.15	4.31	4.95		
	NT2RP2002192	2.9	2.9	7.6	6.65	6.42	5.83		
35	NT2RP2002193	5.21	5.21	5.22	4.76	4.95	5.75		
	NT2RP2002208	5.96	5.96	7.31	4.7	5.67	5.14		
	NT2RP2002219	2.2	2.2	1.8	2.22	3.2	2.26		
40	NT2RP2002231	1.72	1.72	3.1	4.11	2.76	5.76		
	NT2RP2002232	2.59	2.59	5.17	3.93	4.7	6.08		
	NT2RP2002235	5.62	5.62	15.07	16.26	16.18	15.18		
	NT2RP2002239	37.02	37.02	67.99	72.09	67.21	63.77		
45	NT2RP2002252	2.64	2.64	3.66	2.63	2.76	2.94		
	NT2RP2002256	4.62	4.62	15.3	11.37	16.99	12.91		
	NT2RP2002257	7.01	7.01	22.77	18.65	25.09	20.6		
50	NT2RP2002259	1.58	1.58	13.91	9.9	12.15	10.49		
	NT2RP2002264	0.6	0.6	3.14	3.2	3.12	3.92		
	NT2RP2002267	3.66	3.66	8.75	8.95	8.3	11.16		
	NT2RP2002270	4.26	4.26	8.23	16.09	10.47	14.71	*	+
55	NT2RP2002281	2.85	2.85	5.66	8.18	6.48	6.54	*	+
	NT2RP2002288	4.32	4.32	6.6	5.33	5.56	4.23		

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	NT2RP2001818	2.72	2.72	2.45	2.35	3.62	2.66		
	NT2RP2001837	5.13	5.13	13.43	10.29	10.16	12.33		
5	NT2RP2001839	17.02	17.02	-83.84	60.14	71.06	82.26		
	NT2RP2001861	2	2	6.37	3.16	3.52	3.87		
	NT2RP2001869	2.64	2.64	6.54	4.35	5.77	8.84		
	NT2RP2001876	12.15	12.15	27.71	24.54	24.93	23.67		
10	NT2RP2001878	2.32	2.32	3.96	2.95	3.32	4.95		
	NT2RP2001881	3.72	3.72	5.4	9.67	12.64	12.16	**	+
	NT2RP2001883	2.63	2.63	6.8	4.33	5.42	6.35		
15	NT2RP2001884	13.59	13.59	23.56	15.33	10.54	23.6		
	NT2RP2001885	3.27	3.27	5.49	2.88	4.39	4.82		
	NT2RP2001898	10.76	10.76	80.37	69.48	88.43	73.46		
	NT2RP2001900	3.38	3.38	4.03	2.61	4.93	10.26		
20	NT2RP2001903	3.73	3.73	7.71	5.57	5.7	8.2		
	NT2RP2001907	3.1	3.1	8.56	5.05	7.56	6.72		
	NT2RP2001915	2.89	2.89	5.06	4.06	3.08	7.19		
25	NT2RP2001921	4.04	4.04	10.3	13.02	12.45	19.33	*	+
	NT2RP2001926	2.75	2.75	8.25	3.55	5.3	5.64		
	NT2RP2001933	5.65	5.65	52.55	43.62	43.55	48.58		
	NT2RP2001936	1.54	1.54	5.03	2.8	2.96	3.8		
30	NT2RP2001943	25.33	25.33	49.4	47.71	40.48	51.65		
	NT2RP2001946	3.05	3.05	4.3	3.41	4.51	6.1		
	NT2RP2001947	3.18	3.18	3.44	3.93	3.21	6.88		
35	NT2RP2001948	3.59	3.59	10.79	5.71	7.29	19.72		
	NT2RP2001956	5.24	5.24	12.73	11.54	9.42	9.89		
	NT2RP2001969	4.05	4.05	7.82	3.24	5.7	6		
	NT2RP2001976	2.9	2.9	6.39	5.68	6.95	6.41		
40	NT2RP2001978	3.26	3.26	6.08	4.18	4.83	6.03		
	NT2RP2001985	2.14	2.14	3.8	2.56	4.63	2.51		
	NT2RP2001991	3.34	3.34	5.7	1.53	4.62	5.19		
45	NT2RP2001997	3.16	3.16	8.43	5.31	7.47	6.98		
	NT2RP2002015	136.21	136.21	265.98	266.8	340.89	272.43		
	NT2RP2002017	3.24	3.24	6.06	2.06	3.3	2.57		
	NT2RP2002025	6.08	6.08	51.73	31.83	26.94	37.84		
50	NT2RP2002030	6.06	6.06	11.95	9.77	8.07	8.76		
	NT2RP2002032	2.31	2.31	4.95	2.39	3.81	2.55		
	NT2RP2002033	3	3	6.71	3.53	7.57	8.05		
	NT2RP2002041	3.5	3.5	6.37	2.76	3.15	9		
55	NT2RP2002046	3	3	6.88	2.99	7.4	6.01		

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	NT2RP2001526	3.88	3.88	8.49	7.01	5.27	3.83		
	NT2RP2001529	9.87	9.87	53.78	44.74	55.72	60.88		
5	NT2RP2001536	1.63	1.63	-4.17	3.81	4.52	4.71		
	NT2RP2001538	83.44	83.44	178.68	132.75	146.73	155.87		
	NT2RP2001547	4.96	4.96	14.87	16.77	19.21	17.77	*	+
	NT2RP2001560	6.28	6.28	21.64	25.41	28.19	27.75	*	+
10	NT2RP2001562	5.56	5.56	5.57	5.94	6.75	4.64		
	NT2RP2001566	7.96	7.96	9.24	8.22	7.91	8.56		
	NT2RP2001569	4.26	4.26	8.71	6.09	5.65	8.28		
15	NT2RP2001576	3.95	3.95	11.58	13.42	9	12.82		
	NT2RP2001581	47.15	47.15	130.15	121.19	112.28	129.54		
	NT2RP2001597	3.73	3.73	7.88	8.57	8.3	13.3		
	NT2RP2001601	2.37	2.37	4.81	3.67	4.81	3.34		
20	NT2RP2001613	2.74	2.74	4.87	2.83	2.72	4.15		
	NT2RP2001628	3.42	3.42	3.97	3.14	3.04	7.84		
	NT2RP2001634	8.64	8.64	13.94	16.57	23.67	17.67	*	+
25	NT2RP2001635	2.51	2.51	5.92	5.63	5.53	4.72		
	NT2RP2001660	4.27	4.27	16.91	5.9	5.54	10.06		
	NT2RP2001662	1.49	1.49	4.07	4.5	4.44	3.47		
	NT2RP2001663	2.82	2.82	5.09	10.37	8.21	9.74	**	+
30	NT2RP2001672	3.28	3.28	3.82	3.88	4.09	4	*	+
	NT2RP2001675	4.1	4.1	5.01	5.23	4.73	5.06		
	NT2RP2001677	9.58	9.58	18.2	20.9	26.67	19.74	*	+
35	NT2RP2001678	4.84	4.84	6.73	4.6	4.83	4.5		
	NT2RP2001683	1.89	1.89	3.12	4.6	4.72	2.78		
	NT2RP2001699	3.15	3.15	6.16	6.5	5.84	4.88		
	NT2RP2001707	1.24	1.24	3.19	3.42	4.13	4.8	*	+
40	NT2RP2001720	1.47	1.47	3.6	3.91	3	2.72		
	NT2RP2001721	2.26	2.26	4.57	5.53	3.96	3.66		
	NT2RP2001740	12	12	60.21	52.38	79.71	54.73		
45	NT2RP2001748	6.43	6.43	10.8	8.75	10.25	8.55		
	NT2RP2001755	5.51	5.51	4.96	3.71	4.62	2.69	*	-
	NT2RP2001762	1.25	1.25	2.01	3.87	2.56	3.52	*	+
	NT2RP2001768	1.91	1.91	4.7	6.7	5.55	4.55		
50	NT2RP2001769	3.06	3.06	5.86	10.42	5.06	11.86		
	NT2RP2001784	3.62	3.62	6.23	7.06	6.02	6.91		
	NT2RP2001805	2.33	2.33	5.61	6.02	4.93	6.6		
55	NT2RP2001813	2.75	2.75	3.73	1.84	1.98	1.94	*	-
	NT2RP2001817	3.16	3.16	4.49	4.03	5.32	3.45		

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	NT2RP2001277	3.11	3.11	7.02	4.91	6.22	10.82		
	NT2RP2001290	2.71	2.71	6.46	4.42	5.61	6.01		
5	NT2RP2001295	5.46	5.46	9.44	5.13	5.98	7.92		
	NT2RP2001297	118.17	118.17	120.73	139.11	97.16	145.76		
	NT2RP2001301	9.12	9.12	18.56	15.89	19.62	14.24		
10	NT2RP2001312	2.7	2.7	5.68	5.6	4.59	6.04		
	NT2RP2001327	4.73	4.73	5.69	6.39	8.53	11.86		
	NT2RP2001328	8.44	8.44	20.87	16.32	23.25	23.16		
	NT2RP2001341	4.59	4.59	9.22	3.06	7.65	7.21		
15	NT2RP2001347	3.09	3.09	8.54	5.54	9.55	6.9		
	NT2RP2001366	10.33	10.33	48.06	54.83	51.5	52.33		
	NT2RP2001378	2.33	2.33	3.77	3.74	4.64	5.02		
20	NT2RP2001381	2.82	2.82	6.86	5.79	6.62	8.37		
	NT2RP2001388	3.25	3.25	6.71	4.54	5.11	5.2		
	NT2RP2001391	443.52	443.52	734.13	742.83	990.71	747.95		
	NT2RP2001392	2.98	2.98	6.43	4.58	3.16	4.18		
25	NT2RP2001394	3.3	3.3	8.55	8.35	6.09	10.15		
	NT2RP2001397	5.04	5.04	6.79	7.33	5.68	12.2		
	NT2RP2001400	3.1	3.1	6.4	3.43	6.25	2.92		
30	NT2RP2001408	3.31	3.31	6.13	4.02	5.97	5.62		
	NT2RP2001420	5.63	5.63	12.09	8.09	9.97	9.17		
	NT2RP2001423	4.71	4.71	9.71	6.21	8.29	7.19		
	NT2RP2001427	2.68	2.68	5.32	3.69	4.61	5.49		
35	NT2RP2001428	2.71	2.71	7.13	5.49	3.78	3.03		
	NT2RP2001436	4.27	4.27	8.85	5.84	2.85	4.84		
	NT2RP2001440	2.89	2.89	7.34	10.24	10.15	11.98	*	+
	NT2RP2001445	2.43	2.43	6.75	5.86	5.55	5.89		
40	NT2RP2001449	4.37	4.37	6.41	5	4.74	5.02		
	NT2RP2001450	3.19	3.19	6.75	2.26	5.4	8.59		
	NT2RP2001467	4.53	4.53	10.28	5.32	4.72	6.5		
45	NT2RP2001469	4.74	4.74	6.79	8.22	11.04	7.18		
	NT2RP2001480	6.54	6.54	26.68	14.98	12.63	15.42		
	NT2RP2001495	5.86	5.86	11.96	8.16	9.04	10.39		
	NT2RP2001499	8.25	8.25	16.78	10.05	14.46	9.66		
50	NT2RP2001506	2.79	2.79	7.24	5.32	8.19	5.33		
	NT2RP2001508	10.59	10.59	13.66	18.74	20.49	21.92	**	+
	NT2RP2001511	6.41	6.41	9.74	6.08	8.63	6.53		
55	NT2RP2001514	7.04	7.04	7.02	7.24	6.44	6.38		
	NT2RP2001520	2.93	2.93	4.84	2.6	3.19	2.87		

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	NT2RP2000973	0.9	0.9	1.47	2.56	2.64	1.57	*	+
	NT2RP2000985	2.69	2.69	6	9.28	6.29	13.98		
5	NT2RP2000987	1.89	1.89	3.31	4.54	3.17	1.66		
	NT2RP2000997	13.83	13.83	23.99	38.12	29.73	40.96	*	+
	NT2RP2001024	2.86	2.86	5.34	3.61	3.12	2.36		
10	NT2RP2001028	4.66	4.66	4.2	2.65	4.09	0.99		
	NT2RP2001036	5.14	5.14	8.86	6.16	6.44	4.91		
	NT2RP2001039	1.08	1.08	3.18	3.47	1.14	2.24		
	NT2RP2001044	1.13	1.13	2.5	2.53	1.89	2.8		
15	NT2RP2001056	4.97	4.97	28.32	20.16	26.9	18.16		
	NT2RP2001065	2.38	2.38	6.24	7.45	7.4	5.69		
	NT2RP2001067	2.98	2.98	5.38	4.12	5.29	2.41		
20	NT2RP2001070	3.3	3.3	7.63	4.72	6.17	3.58		
	NT2RP2001081	2.91	2.91	8.19	4.8	6.68	3.9		
	NT2RP2001087	3.93	3.93	2.36	2.06	2.92	1.61		
	NT2RP2001094	0.69	0.69	1.37	1.25	1.15	1.04		
25	NT2RP2001119	2.02	2.02	6.11	5.86	4.44	4.35		
	NT2RP2001127	1.53	1.53	4.04	2.69	1.85	2.1		
	NT2RP2001133	2.45	2.45	4.73	4.06	3.61	3.6		
30	NT2RP2001137	2.68	2.68	4.07	2.3	2.82	2.9		
	NT2RP2001142	3.88	3.88	7.47	3.37	2.83	2.42		
	NT2RP2001149	2.7	2.7	2.98	2.11	3.39	1.32		
	NT2RP2001168	6	6	7.81	6.8	7.01	5.75		
35	NT2RP2001173	4.15	4.15	7.88	3.98	3.09	5.44		
	NT2RP2001174	9.23	9.23	14.98	14.12	15.45	18.01		
	NT2RP2001184	2.78	2.78	5.46	4.21	7.18	4.64		
	NT2RP2001196	1.62	1.62	5.93	3.14	3.28	2.94		
40	NT2RP2001200	3.85	3.85	9.36	5.02	4.25	7.88		
	NT2RP2001218	2.29	2.29	5.69	2.55	3.89	3.6		
	NT2RP2001223	2.65	2.65	5.03	1.95	3.69	3.31		
45	NT2RP2001226	4.34	4.34	10.19	6.95	5.72	7.35		
	NT2RP2001227	4.45	4.45	6.12	3.2	3.62	6.01		
	NT2RP2001232	6.44	6.44	13.95	7.13	9.79	13.66		
	NT2RP2001233	4.02	4.02	10.57	7.04	7.77	8.01		
50	NT2RP2001245	4.21	4.21	8.03	9.47	11.82	11.16	*	+
	NT2RP2001246	6.3	6.3	9.84	9	11.28	11.57		
	NT2RP2001268	6.19	6.19	18.1	17.61	16.26	18.55		
55	NT2RP2001270	4.78	4.78	9.11	5.68	8.7	10.04		
	NT2RP2001276	4.92	4.92	13.29	12.73	10.92	12.73		

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	NT2RP2000712	1.43	1.43	5.35	3.39	3.57	2.95		
	NT2RP2000715	3.42	3.42	7.43	6.04	7.56	4.49		
5	NT2RP2000720	4.92	4.92	11.76	7.24	8	7.11		
	NT2RP2000731	3.92	3.92	9.15	3.7	4.5	2.61		
	NT2RP2000739	3.23	3.23	5.67	2.62	3.32	5.65		
	NT2RP2000748	1.59	1.59	4.2	1.42	1.81	1.62		
10	NT2RP2000749	11.84	11.84	21.88	14.4	8.47	13.91		
	NT2RP2000758	1.6	1.6	3.17	2.65	6	1.17		
	NT2RP2000764	1.51	1.51	5.74	2.95	5.22	1.95		
	NT2RP2000766	9.08	9.08	52.24	46.37	59.37	52.89		
15	NT2RP2000777	12.28	12.28	18.43	26.91	28.56	24.47	**	+
	NT2RP2000786	21.32	21.32	73.91	55.85	67.59	58.16		
	NT2RP2000793	5.32	5.32	6.9	4.32	3.57	4.38	*	-
20	NT2RP2000796	5.32	5.32	7.41	7.38	9.17	6.66		
	NT2RP2000809	3.25	3.25	8.3	6.46	4.69	5.45		
	NT2RP2000812	6.65	6.65	17.51	16.43	14.35	16.89		
25	NT2RP2000814	4.16	4.16	4.97	3.75	4.6	3.29		
	NT2RP2000816	1.84	1.84	5.64	4.64	5.19	3.58		
	NT2RP2000818	3.28	3.28	5.19	3.18	3.66	1.95		
	NT2RP2000819	2.76	2.76	5.79	3.03	3.05	1.94		
30	NT2RP2000841	4.35	4.35	4.51	2.17	2.48	1.65	**	-
	NT2RP2000842	7.8	7.8	9.57	13.62	14.25	12.66	**	+
	NT2RP2000845	2.52	2.52	8.31	7.51	6.76	6.93		
35	NT2RP2000863	2.45	2.45	3.48	3.82	3.37	2.47		
	NT2RP2000880	5.96	5.96	11.61	9.5	11.13	10.25		
	NT2RP2000892	4.3	4.3	6.43	6.54	6.97	5.01		
	NT2RP2000894	5.59	5.59	11.88	5.41	5.59	2.16		
40	NT2RP2000903	5.71	5.71	9.12	10.73	11.92	7.44		
	NT2RP2000906	4.56	4.56	5.39	2.63	3.78	2.19	*	-
	NT2RP2000910	4.34	4.34	4.26	2.9	2.7	1.68	**	-
45	NT2RP2000931	10.97	10.97	18.36	20.51	19.28	24.6		
	NT2RP2000932	2.86	2.86	5.43	4.8	4.72	4.21		
	NT2RP2000938	18.41	18.41	42.99	35.71	30.01	43.52		
	NT2RP2000943	7.02	7.02	14.98	18.7	14.88	14.48		
50	NT2RP2000957	3.19	3.19	4.11	4.26	3.66	2.71		
	NT2RP2000958	7	7	6.84	10.43	12.36	7.8		
	NT2RP2000959	9.88	9.88	14.99	13.92	17.38	10.69		
	NT2RP2000965	5.05	5.05	7.82	15.73	18.97	16.02	**	+
55	NT2RP2000970	2.31	2.31	6.72	5.14	5	4.62		

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	NT2RP2000479	3.3	3.3	7.51	5.33	5.71	5.06		
	NT2RP2000498	3.07	3.07	6.25	4.48	5.09	3.9		
5	NT2RP2000503	2.47	2.47	-4.46	2.54	2.82	1.52		
	NT2RP2000510	4.01	4.01	6.19	5.08	6.45	3.7		
	NT2RP2000514	2.65	2.65	2.51	1.94	2.25	1.63	*	-
	NT2RP2000516	4.72	4.72	9.77	4.92	5.29	5.18		
10	NT2RP2000523	2.21	2.21	3.17	1.92	2.44	2.63		
	NT2RP2000533	17.82	17.82	29.05	22.57	27.56	30.78		
	NT2RP2000540	1.98	1.98	4.66	3.01	5.41	5.18		
15	NT2RP2000547	3.1	3.1	5.26	4.38	5.27	3.71		
	NT2RP2000557	4.26	4.26	6.96	4.34	6.5	3.32		
	NT2RP2000558	3.43	3.43	7.17	6.43	7.11	8.26		
	NT2RP2000564	3.04	3.04	7.2	3.49	8.03	4.77		
20	NT2RP2000565	4.54	4.54	11.07	7.64	9.24	9.98		
	NT2RP2000583	14.8	14.8	44.9	49.6	34.93	49.08		
	NT2RP2000591	0.81	0.81	3.81	1.53	2.61	1.21		
25	NT2RP2000599	1.85	1.85	4.1	1.97	3.43	2.36		
	NT2RP2000601	1.78	1.78	4.67	1.28	2.48	1.3		
	NT2RP2000603	2.58	2.58	4.44	2.54	2.84	2.98		
	NT2RP2000610	3.77	3.77	7.23	6.32	7.62	5.53		
30	NT2RP2000614	75.85	75.85	129.42	130.63	184.38	188.58	*	+
	NT2RP2000616	1.81	1.81	4.89	3.9	5.1	3.83		
	NT2RP2000617	2.17	2.17	6.73	5.78	6.82	6.26		
35	NT2RP2000623	3.1	3.1	5.36	3.46	5.1	3.49		
	NT2RP2000634	1.56	1.56	3.92	2.29	3.34	2.02		
	NT2RP2000636	3.78	3.78	8.64	6.27	7.6	6.62		
	NT2RP2000638	4.37	4.37	8.91	4.57	7.41	5.69		
40	NT2RP2000644	2.22	2.22	5.47	3.41	4.16	3.45		
	NT2RP2000649	8.96	8.96	15.76	13.65	17.22	13.07		
	NT2RP2000652	3.35	3.35	4.58	3.57	4.36	2.72		
45	NT2RP2000656	3.73	3.73	6.93	4.83	3.91	4.08		
	NT2RP2000658	1.08	1.08	2.64	1.51	3.18	1.43		
	NT2RP2000663	4.23	4.23	6.9	5.98	7.21	5.9		
	NT2RP2000664	4.24	4.24	10.24	12.72	12.54	16.44	*	+
50	NT2RP2000668	7.49	7.49	26.84	16.92	20.41	17.17		
	NT2RP2000678	1.77	1.77	3.19	1.77	2.09	1.13		
	NT2RP2000694	4.89	4.89	8.39	11.06	13	13.36	**	+
55	NT2RP2000704	1.8	1.8	5.63	2.99	3.13	3.67		
	NT2RP2000710	4.51	4.51	9.96	6.72	8.08	7.23		

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	NT2RP2000240	2.01	2.01	5.25	2.49	3.41	1.45		
	NT2RP2000248	4.29	4.29	6.09	2.82	2.39	0.96	*	-
5	NT2RP2000256	5.7	5.7	8.25	5.62	6.22	5.44		
	NT2RP2000257	3.47	3.47	6.92	4.86	6.52	4.34		
	NT2RP2000258	1.53	1.53	3.83	3.88	2.93	3.02		
	NT2RP2000261	2.95	2.95	3.94	4.47	3.59	2.91		
10	NT2RP2000270	3.12	3.12	6.26	6.66	4.06	4.3		
	NT2RP2000274	1.78	1.78	3.87	3.48	5.16	2.56		
	NT2RP2000277	2.18	2.18	6.13	3.19	4.02	2.98		
15	NT2RP2000279	2.26	2.26	4.92	2.43	2.52	2.17		
	NT2RP2000283	5.75	5.75	27.65	21.53	27.32	24.78		
	NT2RP2000288	6.29	6.29	7.46	8.9	10.89	8.64	*	+
	NT2RP2000289	1.12	1.12	2.79	3.09	2.77	1.93		
20	NT2RP2000297	2.57	2.57	5.7	4.8	4.53	6.09		
	NT2RP2000298	3.61	3.61	9.64	8.51	7.66	8.24		
	NT2RP2000310	1.43	1.43	2.3	2.46	3.41	1.7		
25	NT2RP2000327	2.12	2.12	3.96	3.13	3.49	1.57		
	NT2RP2000328	6.95	6.95	11.56	13.43	16.7	14.68	*	+
	NT2RP2000329	10.73	10.73	10.17	17.55	23.92	18.52	**	+
	NT2RP2000333	6.35	6.35	6.4	6.83	7.17	4.64		
30	NT2RP2000337	2.05	2.05	5.16	4.43	5.32	5.31		
	NT2RP2000346	2.55	2.55	5.18	7.2	5.63	4.95		
	NT2RP2000357	1.57	1.57	6.87	5.48	5.14	5.35		
35	NT2RP2000358	2.09	2.09	4.52	5.03	4.9	4.01		
	NT2RP2000366	3.23	3.23	4.08	4.16	4.25	2.32		
	NT2RP2000369	7.22	7.22	9.94	44.13	45.2	44.34	**	+
	NT2RP2000376	26.92	26.92	108.62	84.48	134.63	85.95		
40	NT2RP2000394	6.49	6.49	5.92	5.08	8.52	4.21		
	NT2RP2000396	2.71	2.71	6.55	7.52	6.8	5.02		
	NT2RP2000412	4.48	4.48	23.45	21.42	24.93	20.49		
45	NT2RP2000414	8.03	8.03	18.69	23.83	18.98	23.37		
	NT2RP2000420	1.12	1.12	4.11	3.54	3.25	1.97		
	NT2RP2000422	6.41	6.41	13.18	17.56	17.88	18.67	*	+
	NT2RP2000426	21.59	21.59	80.94	87.94	110.97	74.98		
50	NT2RP2000428	24.92	24.92	43.91	34.21	35.59	30.95		
	NT2RP2000438	5.06	5.06	5.17	5.62	6.94	5.11		
	NT2RP2000447	4.14	4.14	9.68	7.3	7.08	7.16		
	NT2RP2000448	3.03	3.03	4.63	4.57	3.57	3.17		
55	NT2RP2000459	2.47	2.47	4.93	2.82	3.15	2.09		

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	NT2RP2000056	3.68	3.68	6.23	5.67	6.89	5.8		
	NT2RP2000057	60.79	60.79	174.83	212.63	239.81	221.98	*	+
5	NT2RP2000067	3.1	3.1	3.86	2.98	4.36	5.72		
	NT2RP2000070	2.91	2.91	6.27	5.7	5.95	8.21		
	NT2RP2000076	1.66	1.66	4.45	2.98	3.58	3.23		
	NT2RP2000077	1.67	1.67	4.73	2.43	4.94	3.14		
10	NT2RP2000079	3.76	3.76	9.24	5.15	4.81	5.47		
	NT2RP2000088	2.9	2.9	5.22	2.18	3.07	2.21		
	NT2RP2000091	5.84	5.84	6.54	6.62	8.28	6.72		
15	NT2RP2000092	4.37	4.37	6.7	6.06	7.67	5.65		
	NT2RP2000097	2.74	2.74	3.39	3.4	4.13	4.13	*	+
	NT2RP2000098	3.44	3.44	6.83	6.69	9.01	6.27		
	NT2RP2000108	1.93	1.93	7.24	4.8	6.31	6.68		
20	NT2RP2000114	1.95	1.95	3.65	2.58	4.41	2.9		
	NT2RP2000116	3.17	3.17	7.36	5.35	3.85	9.42		
	NT2RP2000119	3.14	3.14	7.16	4.58	7.96	5.6		
25	NT2RP2000120	3.91	3.91	7.62	5.57	8.5	5.8		
	NT2RP2000126	2.86	2.86	4.86	3.88	5.1	3.44		
	NT2RP2000133	1.83	1.83	3.66	3.13	4.05	2.01		
	NT2RP2000147	6.28	6.28	12.88	11.64	6.51	8.58		
30	NT2RP2000153	4.61	4.61	9.55	10.57	6.49	12.05		
	NT2RP2000156	3.27	3.27	8.24	5.59	6.55	4.07		
	NT2RP2000157	3.7	3.7	6.33	6.57	5.02	4.15		
35	NT2RP2000161	4.45	4.45	8.82	7.52	7.5	6.02		
	NT2RP2000168	4.22	4.22	12.63	3.94	6.03	3.88		
	NT2RP2000173	12.56	12.56	81.37	72.12	92.12	78.68		
	NT2RP2000175	1.9	1.9	3.78	2.03	3.33	4.43		
40	NT2RP2000178	2.06	2.06	6.06	3.23	3.01	3.81		
	NT2RP2000183	1.64	1.64	7.82	5.82	6.21	6.03		
	NT2RP2000195	3.1	3.1	6.65	6	6.71	3.22		
45	NT2RP2000204	73.6	73.6	93.43	102.95	40.16	62.34		
	NT2RP2000205	4	4	6.56	3.91	5.74	4.5		
	NT2RP2000208	3.06	3.06	9.42	4.23	6.77	3.36		
	NT2RP2000224	13.3	13.3	31.75	18.34	21.15	20.93		
50	NT2RP2000230	9.96	9.96	18.99	12.16	16.4	11.95		
	NT2RP2000231	4.3	4.3	7.41	4.24	3.43	4.54		
	NT2RP2000232	1.08	1.08	2.75	1.53	2.45	0.74		
55	NT2RP2000233	8.04	8.04	60.44	47.2	64.72	52.31		
	NT2RP2000239	3	3	4.7	8.93	8.93	7.01	**	+

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	NT2RP1001449	6.23	6.23	7.5	6.29	8.21	4.63		
	NT2RP1001457	4.09	4.09	4.21	2.11	2.26	2.63	**	-
5	NT2RP1001459	21.54	21.54	132.97	90.96	107.97	81.08		
	NT2RP1001466	5.73	5.73	14.97	11.31	9.39	10.99		
	NT2RP1001475	2.45	2.45	6.31	5.98	6.67	3.9		
	NT2RP1001482	3.93	3.93	9.18	15.88	13.03	8.2		
10	NT2RP1001494	1.61	1.61	4.6	4.34	4.18	2.25		
	NT2RP1001500	3.39	3.39	8.13	8.09	8.65	7.42		
	NT2RP1001517	5.11	5.11	7.37	4.41	5.38	2.36		
15	NT2RP1001540	4.74	4.74	5.03	4.6	4.86	3.11		
	NT2RP1001543	1.02	1.02	1.83	1.49	1.12	0.98		
	NT2RP1001546	22.51	22.51	51.51	34.99	22.76	33.42		
	NT2RP1001550	9.33	9.33	21.4	14.35	12.21	13.42		
20	NT2RP1001553	2.07	2.07	6.07	5.69	6.04	4.45		
	NT2RP1001555	36.28	36.28	58.55	41.1	53.63	54.35		
	NT2RP1001563	2.28	2.28	3.44	2.07	2.24	1.31		
25	NT2RP1001569	9.43	9.43	16	17.31	18.21	13.04		
	NT2RP1001584	15.6	15.6	19.66	28.1	32.53	25.83	**	+
	NT2RP1001599	1.18	1.18	1.95	1.27	1.19	1.24		
	NT2RP1001616	5	5	11.95	9.49	6.7	8.95		
30	NT2RP1001654	11.78	11.78	18.07	16.27	16.54	18.38		
	NT2RP1001665	2.77	2.77	4.72	2.73	2.05	2.05		
	NT2RP1001679	76.31	76.31	195.7	199.2	240.87	222.46		
35	NT2RP1001681	10.11	10.11	15.1	20.39	21.03	23.99	**	+
	NT2RP1001694	3.58	3.58	3.82	2.45	2.38	1.97	**	-
	NT2RP2000001	5.23	5.23	5.53	4.17	4.54	3.74	**	-
	NT2RP2000006	3.49	3.49	7.32	4.12	3.88	3.4		
40	NT2RP2000007	3.18	3.18	6.56	4.68	5.66	4.92		
	NT2RP2000008	2.77	2.77	6.72	3.66	5.3	4.9		
	NT2RP2000010	2.89	2.89	5.59	2.99	5.06	2.5		
45	NT2RP2000011	7.08	7.08	17.96	14.55	14.74	15.15		
	NT2RP2000027	2.28	2.28	7.42	4.52	4.89	3.61		
	NT2RP2000028	22.93	22.93	62.54	46.48	51.47	53.47		
	NT2RP2000032	2.5	2.5	5.85	3.11	3.71	6.42		
50	NT2RP2000040	11.57	11.57	23.92	14.38	14.5	23.1		
	NT2RP2000042	5.28	5.28	10.32	6.89	7.21	12.64		
	NT2RP2000045	5.7	5.7	9.42	5.27	6.45	6.3		
55	NT2RP2000051	3.16	3.16	6.29	9.23	9.96	9.53	**	+
	NT2RP2000054	2.55	2.55	6.42	3.81	5.42	2.53		

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	NT2RP1001020	1.87	1.87	4.11	2.08	3.75	2.36		
	NT2RP1001023	62.79	62.79	145.09	101.48	105.86	143.96		
5	NT2RP1001027	18.11	18.11	-82.66	51.63	59.05	68.9		
	NT2RP1001031	1.83	1.83	3.31	2.05	3.53	2.25		
	NT2RP1001033	2.43	2.43	6.09	5.68	5.98	4.27		
	NT2RP1001042	2.94	2.94	6.47	2.99	3.85	2.04		
10	NT2RP1001045	15.95	15.95	23.24	40.66	47.04	44.53	**	+
	NT2RP1001073	6.64	6.64	10.57	8.32	10.46	7.33		
	NT2RP1001079	2.91	2.91	6.37	2.16	2.58	1.48		
15	NT2RP1001080	2.16	2.16	4.89	6.88	4.2	4.56		
	NT2RP1001113	1.07	1.07	3.64	3.55	3.94	3.26		
	NT2RP1001159	21.42	21.42	43.84	22.89	23.31	34.25		
	NT2RP1001173	1.7	1.7	3.07	1.38	4.28	1.52		
20	NT2RP1001176	7.4	7.4	10.13	13	9.31	13.95		
	NT2RP1001177	5.31	5.31	5.75	3.01	5.5	2.02		
	NT2RP1001185	6.42	6.42	9.37	3.79	4.63	2.73	*	-
25	NT2RP1001199	3.9	3.9	7.67	6.93	5.22	3.28		
	NT2RP1001205	7.78	7.78	19.46	16.66	12.64	23.28		
	NT2RP1001215	1.82	1.82	5.02	3.79	4.12	3.15		
	NT2RP1001225	4.54	4.54	7.96	7.56	8.77	6.31		
30	NT2RP1001245	7.27	7.27	10.86	19.68	21.03	22.13	**	+
	NT2RP1001247	2.04	2.04	4.01	1.77	2.89	1.67		
	NT2RP1001248	2.81	2.81	6.79	3.94	4.63	2.4		
35	NT2RP1001253	5.02	5.02	6.39	4.48	4.38	3.32		
	NT2RP1001286	6.18	6.18	7.69	3.79	3.88	4.12	**	-
	NT2RP1001294	2.4	2.4	4.47	3.6	2.73	4.18		
	NT2RP1001302	2.46	2.46	4.51	4.89	2.9	5.39		
40	NT2RP1001310	15.54	15.54	34.01	21.13	20.75	27.15		
	NT2RP1001311	1.9	1.9	3.22	2.66	3.16	2.38		
	NT2RP1001313	2.6	2.6	7.72	5.45	7.85	5.78		
45	NT2RP1001324	2.47	2.47	5.3	3.34	4.17	2.35		
	NT2RP1001349	3.3	3.3	6.29	3.63	3.92	2.14		
	NT2RP1001361	19.41	19.41	18.28	23.28	28.33	24.16	*	+
	NT2RP1001379	3.82	3.82	9.52	4.97	7.97	7.06		
50	NT2RP1001385	2.06	2.06	4.51	4.09	3.89	4.4		
	NT2RP1001395	4.96	4.96	7.86	6.01	6.32	8.13		
	NT2RP1001410	8.75	8.75	20.39	15.74	15.66	9.94		
55	NT2RP1001424	2.39	2.39	3.34	3	3	1.73		
	NT2RP1001432	4.33	4.33	3.86	2.19	1.76	2.05	**	-

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	NT2RP1000769	9.31	9.31	13.98	7.42	8.59	7.19		
	NT2RP1000780	0.86	0.86	1.01	1.87	1.13	0.89		
5	NT2RP1000782	4.25	4.25	12.21	10.24	7.96	9.13		
	NT2RP1000796	3.17	3.17	2.69	4.23	2.99	2.86		
	NT2RP1000797	12.31	12.31	22.78	19.44	18.64	21.78		
	NT2RP1000800	1.13	1.13	3.74	2.46	2.66	1.46		
10	NT2RP1000825	2.38	2.38	2.91	1.04	1.88	0.87	*	-
	NT2RP1000833	2.5	2.5	2.85	0.92	2.39	1.26		
	NT2RP1000834	35.44	35.44	66.57	73.98	90.28	71.45		
15	NT2RP1000836	1.83	1.83	3.43	1.01	2.39	1.04		
	NT2RP1000837	3.36	3.36	6.66	3.22	4.71	3.67		
	NT2RP1000846	1.29	1.29	5.48	1.67	2.84	1.4		
	NT2RP1000847	1.99	1.99	5.49	2.15	5.12	1.64		
20	NT2RP1000851	4.67	4.67	9.32	6.18	7.94	6.72		
	NT2RP1000856	14.31	14.31	17.46	20.38	23.22	19.37	*	+
	NT2RP1000860	2.09	2.09	4.54	4.02	2.74	4.04		
25	NT2RP1000902	5.31	5.31	11.6	6.94	9.91	7.34		
	NT2RP1000903	2.45	2.45	6.26	4.04	3.42	4.24		
	NT2RP1000905	1.76	1.76	4.87	5.36	5.66	10		
	NT2RP1000915	5.51	5.51	10.01	6.72	8.59	9.91		
30	NT2RP1000916	2.31	2.31	5.51	1.78	3.82	2.09		
	NT2RP1000921	9.38	9.38	8.73	8.23	9.13	7.92		
	NT2RP1000943	5.14	5.14	10.76	8.51	8.55	7.2		
35	NT2RP1000944	1.59	1.59	2.21	1.78	1.74	1.15		
	NT2RP1000947	8.5	8.5	14.91	16.51	15.04	14.22		
	NT2RP1000954	2.11	2.11	4.96	2.74	5.55	3.04		
	NT2RP1000958	6.48	6.48	14.73	4.54	10.17	10.21		
40	NT2RP1000959	124.81	124.81	209.45	128.43	72.65	206.1		
	NT2RP1000966	9.96	9.96	12.96	14.28	15.36	21.39		
	NT2RP1000974	2.46	2.46	5.38	3.98	6.08	3.71		
45	NT2RP1000980	3.07	3.07	5.5	4.04	4.53	4.02		
	NT2RP1000981	4.3	4.3	8.09	5.68	7.26	5.27		
	NT2RP1000988	6.45	6.45	10.46	9.62	6.44	7.87		
	NT2RP1001002	2.8	2.8	7.36	3.94	4.57	4.3		
50	NT2RP1001004	4.72	4.72	8.25	3.65	4.9	5.37		
	NT2RP1001007	1.42	1.42	3.42	1.69	3.84	2.03		
	NT2RP1001011	1.94	1.94	5.93	3.82	5.46	4.83		
55	NT2RP1001013	4.45	4.45	9.41	5.92	8.62	5.04		
	NT2RP1001014	2.21	2.21	5.89	3.76	6.64	3.49		

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	NT2RP1000544	1.53	1.53	2.45	1.38	1.24	1.44		
	NT2RP1000547	0.88	0.88	2.45	2	1.63	1.23		
5	NT2RP1000551	1.7	1.7	-2.62	2.13	3.2	1.1		
	NT2RP1000567	1.66	1.66	4.29	2.54	4.29	1.77		
	NT2RP1000574	1.99	1.99	4.28	1.5	3.43	1.38		
10	NT2RP1000577	3.14	3.14	6.01	3.16	5.31	2.05		
	NT2RP1000579	4.64	4.64	6.24	3.27	3.97	2.04		
	NT2RP1000581	5.22	5.22	3.58	2.07	1.61	0.93	**	-
	NT2RP1000593	1.74	1.74	4.39	2.48	3.28	2.3		
15	NT2RP1000604	3.85	3.85	7.75	17.25	13.78	16.39	**	+
	NT2RP1000609	1.15	1.15	2.21	2.84	2.61	1.55		
	NT2RP1000613	1.12	1.12	2.56	1.82	4.29	0.82		
20	NT2RP1000622	5.94	5.94	15.9	14.91	19.42	15.46		
	NT2RP1000627	9.18	9.18	18.96	23.88	21.9	14.86		
	NT2RP1000629	4.18	4.18	5.9	5.92	5.32	3.17		
	NT2RP1000630	6.54	6.54	7.84	7.21	7.67	7.92		
25	NT2RP1000639	0.64	0.64	0.31	1.53	2.04	0.28		
	NT2RP1000640	130.14	130.14	307.77	227.5	176.05	232.29		
	NT2RP1000646	4.14	4.14	9.59	10.19	11.87	12.15	*	+
	NT2RP1000659	2.65	2.65	7	8.91	7.99	6.04		
30	NT2RP1000674	13.48	13.48	28.08	43.62	45.82	56.95	**	+
	NT2RP1000677	3.9	3.9	10.76	11.84	10.19	9.87		
	NT2RP1000679	2.38	2.38	3.76	2.3	2.35	1.05		
35	NT2RP1000688	4.72	4.72	3.34	2.76	2.73	1.83	*	-
	NT2RP1000689	1.44	1.44	1.86	2.03	1.22	1.13		
	NT2RP1000695	1.11	1.11	2.5	2.09	2.44	1.52		
	NT2RP1000701	0.89	0.89	1.08	2.62	2.74	1.71	*	+
40	NT2RP1000702	1.12	1.12	2.28	3.74	4.07	3.15	*	+
	NT2RP1000713	2.29	2.29	2.79	2.8	3.56	2.38		
	NT2RP1000721	4.14	4.14	4.49	4.48	3.92	3.78		
45	NT2RP1000730	3.5	3.5	4.83	2.61	4.5	2.41		
	NT2RP1000733	6.08	6.08	6.56	4.91	8.12	5.65		
	NT2RP1000738	3.18	3.18	8.04	5.16	5.71	7.11		
	NT2RP1000739	1.11	1.11	2.65	4.02	3.09	2.86		
50	NT2RP1000740	1.41	1.41	3.13	3.63	3.57	3.77	*	+
	NT2RP1000746	1.15	1.15	3.58	2.28	3.74	1.37		
	NT2RP1000750	4	4	8.31	10.25	10.72	9.39	*	+
55	NT2RP1000751	33.15	33.15	59.65	67.84	64.22	66.55		
	NT2RP1000767	3.8	3.8	3.64	1.7	2.62	0.62	*	-

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	NT2RP1000326	4.01	4.01	7.67	3.82	4.56	7.85		
	NT2RP1000331	12.16	12.16	24.08	12.19	10.5	20.1		
5	NT2RP1000333	4.18	4.18	-7.52	6.66	6.53	6.98		
	NT2RP1000336	1.45	1.45	4.45	1.35	3.76	1.78		
	NT2RP1000347	3.05	3.05	8.75	7.26	8.31	6.38		
	NT2RP1000348	2.11	2.11	4.14	2.76	3	2.75		
10	NT2RP1000349	2.12	2.12	3.92	3	4.39	4.11		
	NT2RP1000353	40.87	40.87	83.5	51.49	47.8	66.02		
	NT2RP1000356	39.53	39.53	93.37	50.3	56.48	74.42		
15	NT2RP1000357	3.89	3.89	9.63	8.43	8.7	8.72		
	NT2RP1000358	2.85	2.85	6.11	4.23	3.04	5.09		
	NT2RP1000360	11.04	11.04	19.39	12.08	18.42	19.44		
	NT2RP1000363	13.09	13.09	15.39	13.13	13.38	10.01		
20	NT2RP1000376	1.81	1.81	3.8	2.09	1.9	2.24		
	NT2RP1000386	118	118	191.31	146.98	187.97	155.47		
	NT2RP1000407	0.72	0.72	3.16	0.58	0.89	1.2		
25	NT2RP1000409	2.05	2.05	5.39	2.84	6.59	3.83		
	NT2RP1000413	4.78	4.78	8.03	5.86	8.89	10.19		
	NT2RP1000416	1.5	1.5	2.01	0.93	3.17	0.7		
	NT2RP1000418	2.27	2.27	6.69	5.08	6.67	4.85		
30	NT2RP1000420	1.77	1.77	5.19	7.32	7.64	3.7		
	NT2RP1000434	1.48	1.48	4.39	1.27	3.12	1		
	NT2RP1000439	5.02	5.02	9.31	20.62	28.73	24.75	**	+
35	NT2RP1000443	1.8	1.8	3.46	2.24	1.61	1.63		
	NT2RP1000447	2.21	2.21	5.57	2.49	2.87	3.1		
	NT2RP1000448	1.39	1.39	3.58	3.09	4.4	1.41		
	NT2RP1000451	4.2	4.2	6.37	5.72	7.27	7.04		
40	NT2RP1000458	15.1	15.1	10.53	19.73	8.72	23.03		
	NT2RP1000460	7.55	7.55	13.82	8.76	11.49	8.62		
	NT2RP1000465	4.58	4.58	20.97	20.41	19.98	22.46		
45	NT2RP1000468	3.25	3.25	4.64	3.82	4.1	4.45		
	NT2RP1000470	2.38	2.38	5.67	3.99	2.35	3.8		
	NT2RP1000477	1.11	1.11	3.81	1.1	0.84	0.83		
	NT2RP1000478	4.53	4.53	12.55	19.87	18.75	20.39	*	+
50	NT2RP1000481	1.23	1.23	3.89	2.48	4.09	1.2		
	NT2RP1000493	2.44	2.44	3.8	1.74	3.83	0.87		
	NT2RP1000513	13.07	13.07	16.37	17.06	17.57	18.97	*	+
55	NT2RP1000522	6.13	6.13	12.69	13.13	13.08	10.32		
	NT2RP1000533	3.72	3.72	6.17	2.92	4.49	2.17		

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	NT2RP1000068	1.89	1.89	3.99	3.12	3.33	2.43		
	NT2RP1000072	22.9	22.9	74.07	82.91	66.26	95.85		
5	NT2RP1000073	2.18	2.18	2.45	2.68	3.69	3.86	*	+
	NT2RP1000078	2.72	2.72	3.17	2.93	2.3	3.13		
	NT2RP1000079	4.13	4.13	5.32	3.6	4.48	2.5		
10	NT2RP1000080	4.99	4.99	8.13	9.46	12.46	9.46	*	+
	NT2RP1000086	4.15	4.15	3.63	1.31	2.1	3.75		
	NT2RP1000087	1.3	1.3	4.36	3.51	3.21	3.45		
	NT2RP1000089	4.5	4.5	9.98	12.69	11.3	14.93	*	+
15	NT2RP1000090	45.76	45.76	96.6	94.37	53.44	93.42		
	NT2RP1000100	2.17	2.17	4.05	5.23	4.13	3.66		
	NT2RP1000101	3.44	3.44	5.22	4.41	2.88	4.81		
20	NT2RP1000111	3.24	3.24	5.56	4.51	3.9	3.69		
	NT2RP1000112	3.29	3.29	4.08	1.85	3.33	3		
	NT2RP1000124	5.57	5.57	4.96	3.11	5.73	5.5		
	NT2RP1000125	7.28	7.28	19.39	13.69	10.68	16.86		
25	NT2RP1000129	1.81	1.81	4.35	5.14	3.91	4.27		
	NT2RP1000130	2.31	2.31	4.11	5.31	5.62	16.86		
	NT2RP1000154	7.5	7.5	15.63	17.16	12.72	16.37		
30	NT2RP1000163	2.42	2.42	3.51	2.72	2.99	3.59		
	NT2RP1000170	3.42	3.42	4.2	4.96	5.17	5.85	*	+
	NT2RP1000174	3.5	3.5	3.42	1.3	2.38	2.12	**	-
	NT2RP1000181	6.14	6.14	7.22	10.97	14.98	9.38	*	+
35	NT2RP1000191	1.08	1.08	5.61	4.94	3.59	5.71		
	NT2RP1000202	1.06	1.06	1.66	2.02	1.2	2.24		
	NT2RP1000239	1.53	1.53	4.1	2.15	0.94	2.07		
40	NT2RP1000243	2.37	2.37	2.04	1.31	1.14	1.64	**	-
	NT2RP1000255	1.94	1.94	3.02	2.11	2.26	1.78		
	NT2RP1000259	5.27	5.27	9.55	5.53	6.33	4.29		
	NT2RP1000261	2.76	2.76	4.4	2.07	1.64	2.64		
45	NT2RP1000269	5.16	5.16	5.01	7.7	10.51	7.39	*	+
	NT2RP1000271	7.79	7.79	15.88	13.16	15.11	18.48		
	NT2RP1000272	7.71	7.71	13.07	10.72	11.74	11.3		
	NT2RP1000279	2.19	2.19	5.24	2.23	3.91	2.62		
50	NT2RP1000290	6.61	6.61	9.02	12.65	13.52	9.92	*	+
	NT2RP1000293	6.86	6.86	10.91	9.75	8.45	10.92		
	NT2RP1000300	12.42	12.42	11.93	9.96	11.37	10.2	*	-
55	NT2RP1000324	5.16	5.16	6	4.69	5.92	6.97		
	NT2RP1000325	54.42	54.42	101.4	70.46	57.52	78.6		

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	NT2RM4002493	1.35	1.35	3.22	1.96	3.51	2.73		
	NT2RM4002499	34.96	34.96	72.9	59.42	52.07	54.3		
5	NT2RM4002504	5.15	5.15	10.68	10.57	13.51	9.8		
	NT2RM4002506	4.77	4.77	9.4	4.93	7.59	8.53		
	NT2RM4002510	2.03	2.03	3.27	1.66	2.97	2.48		
	NT2RM4002527	1.57	1.57	3.14	1.83	2.31	4.47		
10	NT2RM4002532	2.45	2.45	7.75	5.88	3.37	5.19		
	NT2RM4002534	1.79	1.79	4.8	2.1	3.45	2.94		
	NT2RM4002535	2.5	2.5	6.51	5.89	8.1	6.37		
15	NT2RM4002554	3.29	3.29	5.31	3.31	5.47	3.12		
	NT2RM4002558	6.91	6.91	32.57	32.58	41.54	25.61		
	NT2RM4002565	5.38	5.38	13.6	8.22	9.85	10.53		
	NT2RM4002567	3.34	3.34	5.43	4.21	4.49	7.22		
20	NT2RM4002571	4.48	4.48	15.61	11.95	16.47	15.24		
	NT2RM4002572	5.57	5.57	17.2	13.7	9.59	13.48		
	NT2RM4002577	7.76	7.76	15.25	6.59	5.87	5.65		
25	NT2RM4002583	1.08	1.08	3.58	2.28	4.83	2.44		
	NT2RM4002584	1.64	1.64	5.67	3.24	5.74	5.56		
	NT2RM4002593	3.29	3.29	5.17	2.75	3.91	4.61		
	NT2RM4002594	11.26	11.26	46.5	38.21	55.32	43.58		
30	NT2RM4002604	4.83	4.83	4.64	1.77	2.03	2.89	**	-
	NT2RM4002614	3.48	3.48	3.48	2.52	3.66	2.81		
	NT2RM4002616	1.07	1.07	2.73	2.88	2.71	2.38		
35	NT2RM4002623	1.39	1.39	4.89	3.92	3.72	5.06		
	NT2RM4002634	1.41	1.41	4.38	3.42	4.91	2.96		
	NT2RM4002636	2.22	2.22	3.93	3.92	4.18	4.12		
40	NT2RP1000002	8.82	8.82	52.94	75.1	92.89	81.45	*	+
	NT2RP1000006	4.68	4.68	6.28	4.25	4.48	2.56		
	NT2RP1000015	4.86	4.86	5.27	2.74	1.99	2.28	**	-
	NT2RP1000018	5.45	5.45	5	5.55	4.83	4.96		
45	NT2RP1000034	18.22	18.22	49.95	38.04	30.76	50.07		
	NT2RP1000035	1.93	1.93	3.2	5.26	3.23	3.96		
	NT2RP1000040	1.77	1.77	3.33	2.93	3.28	4.28		
	NT2RP1000042	1.3	1.3	3.44	1.99	3.22	2.38		
50	NT2RP1000048	3.6	3.6	10.24	7.25	9.9	9		
	NT2RP1000050	2.21	2.21	4.71	2.89	4	3.57		
	NT2RP1000056	4.03	4.03	3.74	1.09	0.61	1.96	**	-
55	NT2RP1000058	3.49	3.49	2.03	1.84	2.07	2.48		
	NT2RP1000063	1.77	1.77	3.65	4.09	4	3.83		

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	NT2RM4002251	2.11	2.11	6.2	7.87	5.48	5.17		
	NT2RM4002256	4.38	4.38	10.68	10.7	9.46	9.64		
5	NT2RM4002262	2.85	2.85	6.25	3.34	4.43	9.66		
	NT2RM4002266	3.93	3.93	4.76	2.76	3.55	4.47		
	NT2RM4002276	11.23	11.23	15.55	16.5	28.25	20.64		
	NT2RM4002278	1.89	1.89	4.59	4.33	3.99	5.11		
10	NT2RM4002281	17.71	17.71	59.08	62.68	51.19	59.89		
	NT2RM4002287	2.08	2.08	3.84	2.46	4.21	3.32		
	NT2RM4002294	3.19	3.19	6.99	6.28	6.09	8.69		
15	NT2RM4002298	18.59	18.59	60.14	86.09	89.9	88.75	*	+
	NT2RM4002301	3.2	3.2	6.85	4.63	5.94	4.02		
	NT2RM4002306	4.71	4.71	8.24	4.99	5.31	4.2		
	NT2RM4002323	3.9	3.9	4.06	4.11	4.39	3.11		
20	NT2RM4002334	11.54	11.54	20.76	17.92	20.72	16.95		
	NT2RM4002339	1.78	1.78	3.52	1.33	1.3	1.38		
	NT2RM4002344	2.36	2.36	5.74	2.87	3.57	7.92		
25	NT2RM4002345	3.56	3.56	10.59	5.06	4.63	7.5		
	NT2RM4002352	2.04	2.04	7.67	3.99	5.14	3.74		
	NT2RM4002362	20.38	20.38	24.92	11.23	14.32	15.17	*	-
	NT2RM4002373	2.1	2.1	3.96	3.21	2.55	3.63		
30	NT2RM4002374	2.28	2.28	4.39	2.29	3.58	4.3		
	NT2RM4002376	4.02	4.02	6.03	3.31	2.97	5.52		
	NT2RM4002383	2.8	2.8	8.49	4.76	5.79	4.28		
35	NT2RM4002390	3.03	3.03	6.01	4.06	5.27	7.37		
	NT2RM4002398	5.16	5.16	43.18	33.97	50.73	30.41		
	NT2RM4002409	2.11	2.11	5.93	3.37	4.29	1.9		
	NT2RM4002414	4.73	4.73	6.21	7.37	9.12	14.53		
40	NT2RM4002438	2.07	2.07	5.28	3.03	4.38	7.18		
	NT2RM4002440	2.99	2.99	6.92	5.78	5.32	9.49		
	NT2RM4002446	2.23	2.23	6.08	2.95	4.45	5.7		
45	NT2RM4002450	3.36	3.36	10.01	6.15	7.75	7.24		
	NT2RM4002452	2.13	2.13	6.3	3.67	5.15	7.23		
	NT2RM4002457	2.68	2.68	4.44	2.66	3.26	4.52		
	NT2RM4002458	3.06	3.06	5.77	3.32	5.34	4.04		
50	NT2RM4002460	2.43	2.43	3.68	1.57	2.45	1.43		
	NT2RM4002464	5.4	5.4	12.62	14.39	13.72	14.3		
	NT2RM4002479	4.66	4.66	6.69	4.91	7.98	11.54		
55	NT2RM4002482	4.26	4.26	16.18	10.19	11.5	12.2		
	NT2RM4002489	6.74	6.74	16.91	8.79	5.81	11.68		

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	NT2RM4002054	4.54	4.54	6.97	2.56	4.3	3.89		
	NT2RM4002055	13.72	13.72	74.75	60.51	91.27	61.53		
5	NT2RM4002059	23.73	23.73	31.85	48.05	63.09	52.61	**	+
	NT2RM4002061	3.72	3.72	5.32	3.59	3.69	4.81		
	NT2RM4002062	1.9	1.9	5.41	3.66	2.84	4.26		
	NT2RM4002063	2.21	2.21	8.1	7.64	7.35	3.79		
10	NT2RM4002066	2.07	2.07	5.29	4.42	6.32	4.07		
	NT2RM4002067	2.51	2.51	4.27	3.07	5.19	4.41		
	NT2RM4002073	3.73	3.73	7.24	5.51	7.69	5.16		
15	NT2RM4002074	5.19	5.19	7.35	5.67	7.47	4.49		
	NT2RM4002075	5.13	5.13	5.9	3.16	3.18	2.91	**	-
	NT2RM4002076	3.13	3.13	3.05	1.94	2.52	1.71	*	-
	NT2RM4002078	10.3	10.3	28.06	23.95	20.81	26.64		
20	NT2RM4002081	10.47	10.47	30.87	19.18	17.8	18.22		
	NT2RM4002082	1.25	1.25	3.02	3.85	2.58	1.23		
	NT2RM4002093	2.82	2.82	3.9	4.79	4.66	4.79	*	+
25	NT2RM4002109	4.42	4.42	11.51	13.95	15.12	15.21	*	+
	NT2RM4002115	2.86	2.86	4.51	4.81	4.8	2.52		
	NT2RM4002118	4.48	4.48	6.14	4.3	4.86	4.27		
	NT2RM4002128	3.78	3.78	4.57	2.84	3.31	3.13	*	-
30	NT2RM4002137	3.96	3.96	8.14	10.27	7.51	8.92		
	NT2RM4002139	3.78	3.78	8.98	7.03	7.84	7.87		
	NT2RM4002140	4.04	4.04	9.45	8.87	7.81	10.17		
35	NT2RM4002145	5.99	5.99	17.51	25.81	31.07	24.47	*	+
	NT2RM4002146	4.51	4.51	8.23	8.56	9	8.92		
	NT2RM4002161	2.33	2.33	4.97	1.38	3.15	5.3		
	NT2RM4002174	4.86	4.86	8.02	3.12	4.53	6.15		
40	NT2RM4002178	7.3	7.3	24.43	28.61	33.13	29.27	*	+
	NT2RM4002180	3.47	3.47	11.93	9.27	10.02	11.28		
	NT2RM4002185	5.94	5.94	35.51	31.59	32.34	31.69		
45	NT2RM4002189	1.6	1.6	3.24	3.68	5.59	4.91	*	+
	NT2RM4002194	9.3	9.3	25.94	37.2	29.64	38.23	*	+
	NT2RM4002198	6.09	6.09	7.61	9.37	8.4	10.04	*	+
	NT2RM4002205	4.01	4.01	9.05	6.76	7.86	8.76		
50	NT2RM4002213	5.36	5.36	8.79	8.05	11.99	14.41		
	NT2RM4002216	7.35	7.35	12.58	16.58	23.93	18.16	*	+
	NT2RM4002226	3.84	3.84	9.71	20.85	16.65	16.5	**	+
55	NT2RM4002237	4.19	4.19	10.13	10.37	7.64	13.22		
	NT2RM4002240	1.96	1.96	3.64	3.73	3.71	7.59		

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	NT2RM4001836	3.27	3.27	8.32	3.65	3.54	5.58		
	NT2RM4001841	7.94	7.94	15.82	20.15	20.96	23.33	*	+
5	NT2RM4001842	2.1	2.1	4.44	3.5	3.35	4.85		
	NT2RM4001843	5.65	5.65	14.54	13.34	12.25	14.94		
	NT2RM4001856	4.42	4.42	7.16	7.65	4.71	16.83		
	NT2RM4001858	5.91	5.91	15.86	16.09	17.03	16.93		
10	NT2RM4001861	2.91	2.91	9.57	6.31	8.66	9.28		
	NT2RM4001863	8.06	8.06	9.5	15.16	15.68	11.77	*	+
	NT2RM4001865	5.04	5.04	11.25	7.44	10.24	9.03		
15	NT2RM4001869	5.1	5.1	5.96	5.22	8.45	21.88		
	NT2RM4001873	9.62	9.62	18.43	13.33	15.49	19.21		
	NT2RM4001876	2.24	2.24	6.94	3.65	4.39	7.25		
	NT2RM4001880	3.6	3.6	8.57	5.13	5.41	7.67		
20	NT2RM4001885	5.71	5.71	11.11	7.11	6.56	11.98		
	NT2RM4001889	10.25	10.25	18.24	16.31	15.85	21.33		
	NT2RM4001894	2.61	2.61	6.07	3.58	3.65	3.49		
25	NT2RM4001897	7.87	7.87	20.24	18.41	20.4	23.46		
	NT2RM4001899	3.36	3.36	7.43	4.92	8.19	8.54		
	NT2RM4001905	3	3	4.84	3.3	4.53	7.1		
	NT2RM4001922	2.55	2.55	6.05	3.97	4.84	5.11		
30	NT2RM4001930	2.64	2.64	8.9	2.88	6.53	6.38		
	NT2RM4001938	2.65	2.65	4.91	5.09	5.65	6.43		
	NT2RM4001940	2.73	2.73	6.17	5.91	4.46	5.48		
35	NT2RM4001942	37.36	37.36	32.02	53.86	59.28	82.77	*	+
	NT2RM4001953	4.65	4.65	9.68	5.04	6.79	4.91		
	NT2RM4001965	4.96	4.96	8.82	10.18	8.54	8.39		
	NT2RM4001966	3	3	5.14	6.3	7.45	8.19	*	+
40	NT2RM4001969	2.22	2.22	7.29	5.01	2.95	4.5		
	NT2RM4001974	1.19	1.19	4.61	1.89	2.96	4.83		
	NT2RM4001979	2.09	2.09	6.37	3.39	4.65	7.36		
45	NT2RM4001980	4.3	4.3	7.59	7.58	8.02	9.33		
	NT2RM4001984	2.31	2.31	5.36	2.68	3.49	4.57		
	NT2RM4001987	3.36	3.36	9.66	2.92	4.6	5.01		
	NT2RM4002013	6.62	6.62	15.13	13.47	17.16	19.8		
50	NT2RM4002018	2.31	2.31	5.15	4.09	5.53	7.1		
	NT2RM4002033	3.19	3.19	8.16	4.91	3.27	5.93		
	NT2RM4002034	1.89	1.89	6.19	4.82	4.38	4.03		
55	NT2RM4002044	7.71	7.71	17.9	18.75	12.3	18.5		
	NT2RM4002047	3.88	3.88	5.19	2.68	5.38	9.2		

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	NT2RM4001611	3.31	3.31	4.56	2.33	2.32	2.02	*	-
	NT2RM4001618	7.05	7.05	7.95	6.68	8.98	12.95		
5	NT2RM4001622	13.53	13.53	-19.88	14.67	24.67	28.46		
	NT2RM4001624	1.6	1.6	3.02	3.92	2.66	4.43		
	NT2RM4001625	4.89	4.89	39.6	41.63	47.1	46.46		
	NT2RM4001629	3.82	3.82	8.82	12.09	12.08	13.38	*	+
10	NT2RM4001632	15.28	15.28	24.55	31.07	26.16	25.6		
	NT2RM4001642	3.29	3.29	4.17	2.89	3.62	2.1		
	NT2RM4001647	4.44	4.44	6.83	4.04	5.48	4.67		
15	NT2RM4001650	4.96	4.96	4.94	2.66	2.87	3.79	**	-
	NT2RM4001662	2.18	2.18	5.47	8.31	6.54	9.39	*	+
	NT2RM4001666	2.28	2.28	6.5	6.24	6.17	8.14		
	NT2RM4001670	3.52	3.52	10.77	11.16	10.82	14.91		
20	NT2RM4001682	12.66	12.66	31.6	33.03	26.04	37.07		
	NT2RM4001710	6.7	6.7	38.5	40.58	58.41	40.31		
	NT2RM4001712	4.06	4.06	7.61	10.19	10.7	9.98	*	+
25	NT2RM4001714	10.88	10.88	19.37	18.67	19.3	17.65		
	NT2RM4001715	10.77	10.77	11.6	13.55	16.86	12.99		
	NT2RM4001727	3.41	3.41	5.92	4.83	5.89	7.6		
30	NT2RM4001731	2.6	2.6	10.72	13.46	11.23	11.73		
	NT2RM4001735	12.84	12.84	21.53	22.01	20.88	34.93		
	NT2RM4001739	2.46	2.46	7.3	8.13	5.17	7.14		
	NT2RM4001741	14.41	14.41	29.88	26.98	27.21	32.35		
35	NT2RM4001746	3.65	3.65	6.76	6.89	6.5	5.33		
	NT2RM4001754	3.16	3.16	4.17	3.39	3.62	3.84		
	NT2RM4001757	5.02	5.02	5.78	4.7	6.31	7.97		
	NT2RM4001758	1	1	0.76	1.98	0.65	1.46		
40	NT2RM4001768	4.83	4.83	10.19	8.48	6.91	7.83		
	NT2RM4001775	3.23	3.23	2.76	1.9	1.85	1.71	**	-
	NT2RM4001776	2.56	2.56	4.77	2.47	2.68	2.69		
45	NT2RM4001783	2.88	2.88	3.22	3.12	3.48	3.68		
	NT2RM4001793	4.67	4.67	11.44	12.02	9.6	10.75		
	NT2RM4001810	3.31	3.31	4.46	3.33	3.63	3.11		
	NT2RM4001813	3.9	3.9	4.15	4.71	4.19	5.36		
50	NT2RM4001818	4.06	4.06	11.34	10.43	8.67	10.53		
	NT2RM4001819	2.35	2.35	5.6	2.37	3.02	4.58		
	NT2RM4001823	1.76	1.76	4.48	2.47	4.04	4.27		
55	NT2RM4001828	5.01	5.01	11.49	5.67	7.54	7.51		
	NT2RM4001835	9.75	9.75	18.65	21.12	16.5	26.55		

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	NT2RM4001371	4.71	4.71	8.57	6.58	13.16	12.93		
	NT2RM4001377	8.01	8.01	13.74	8.66	9.23	10.58		
5	NT2RM4001382	17.31	17.31	56.74	35.57	28.32	38.22		
	NT2RM4001384	2.17	2.17	4.11	2.35	3.34	3.75		
	NT2RM4001400	3.78	3.78	5.76	4.87	4.66	5.07		
	NT2RM4001409	2.55	2.55	6.82	4.82	4.63	5.27		
10	NT2RM4001410	5.48	5.48	21.69	17.34	21.09	18.91		
	NT2RM4001411	2.66	2.66	6.5	5.86	6.64	6.77		
	NT2RM4001412	1.75	1.75	3.33	2.39	2.52	2.79		
15	NT2RM4001414	2.18	2.18	4.83	2.45	2.4	3.77		
	NT2RM4001436	8.35	8.35	16.65	10.45	10.13	15.55		
	NT2RM4001437	2.77	2.77	8.85	8.31	11.26	7.67		
	NT2RM4001444	11.57	11.57	25.93	21.1	18.36	18.42		
20	NT2RM4001454	3.62	3.62	7.89	6.16	5.03	4.79		
	NT2RM4001455	5.85	5.85	22.19	18.03	25.08	26.16		
	NT2RM4001483	4.37	4.37	8.77	4.97	5.43	6.46		
25	NT2RM4001489	3.12	3.12	7.04	3.64	4.19	5.46		
	NT2RM4001495	1.35	1.35	5.13	2.85	3.29	3.37		
	NT2RM4001499	1.74	1.74	4.98	3.65	2.77	3.3		
	NT2RM4001515	0.95	0.95	3.38	2.65	5.38	3.41		
30	NT2RM4001519	1.86	1.86	4.6	2.24	4.38	2.85		
	NT2RM4001522	3.8	3.8	7.66	5.36	7.42	9.13		
	NT2RM4001523	5.46	5.46	9.11	4.82	7.58	5.79		
35	NT2RM4001550	11.9	11.9	17.38	16.49	16.76	15.01		
	NT2RM4001553	7.88	7.88	13.4	23.26	23.53	23.87	**	+
	NT2RM4001554	0.86	0.86	1.74	1.85	1.11	2.2		
	NT2RM4001557	2.5	2.5	6.33	5.05	3.36	4.89		
40	NT2RM4001565	1.87	1.87	4.05	2.46	3.34	3.23		
	NT2RM4001566	3.23	3.23	8.57	8.91	10.49	11.42		
	NT2RM4001569	1.47	1.47	5.4	3.35	4.15	1.56		
45	NT2RM4001579	6.57	6.57	16.69	19.23	23.83	18.22		
	NT2RM4001582	4.06	4.06	5.97	2.16	3.17	2.67	*	-
	NT2RM4001589	21.51	21.51	37.16	42.45	55.76	47.57	*	+
	NT2RM4001592	1.37	1.37	2.96	3.02	2.14	3.71		
50	NT2RM4001594	1.98	1.98	4.09	5.4	5.24	5.67	*	+
	NT2RM4001597	2.65	2.65	5.64	5.17	4.97	4.33		
	NT2RM4001605	2.7	2.7	6.18	5.71	5.92	5.93		
55	NT2RM4001609	23.65	23.65	45	61.08	78.89	77.31	*	+
	NT2RM4001610	48.1	48.1	69.16	132.54	132.39	115.22	**	+

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	NT2RM4001084	8.04	8.04	6.52	9.49	10.53	9.53	*	+
	NT2RM4001092	12.61	12.61	109.97	76.93	98.78	73.14		
5	NT2RM4001100	6.72	6.72	20.93	22.35	15.18	18.98		
	NT2RM4001116	1.17	1.17	2.5	2.77	2.96	2.16		
	NT2RM4001119	1.74	1.74	4.82	4.72	4.22	4.15		
	NT2RM4001140	2.65	2.65	7.49	8.27	7.78	9.03		
10	NT2RM4001148	6.59	6.59	13.68	12.1	12.92	15.83		
	NT2RM4001151	3.7	3.7	4.31	1.94	3.65	1.68		
	NT2RM4001155	5.51	5.51	6.36	3.55	5.37	4.63		
15	NT2RM4001157	1.68	1.68	4.95	4.86	2.81	3.82		
	NT2RM4001160	1.57	1.57	3.39	2.99	1.9	2.41		
	NT2RM4001163	42.35	42.35	86.27	106.63	52.13	98.56		
	NT2RM4001187	2.64	2.64	7.25	3.96	4.87	5.2		
20	NT2RM4001191	3.2	3.2	9.95	8.59	6.36	8.72		
	NT2RM4001200	3.83	3.83	6.35	5.52	4.41	4.11		
	NT2RM4001203	9.93	9.93	20.29	22.65	25.36	21.82		
25	NT2RM4001204	3.23	3.23	3.59	3.17	2.83	2.6		
	NT2RM4001217	4.62	4.62	12.16	11.26	13.39	13.72		
	NT2RM4001245	7.31	7.31	17.14	13.76	14.49	17.16		
	NT2RM4001247	3.23	3.23	9.1	5.73	6.57	5.03		
30	NT2RM4001256	2.51	2.51	6.39	3.57	5.48	4.14		
	NT2RM4001258	8.2	8.2	24.68	25.02	24.16	22.89		
	NT2RM4001267	3.43	3.43	4.83	3.93	4.3	8.08		
	NT2RM4001273	4.23	4.23	8.38	7.39	6.96	8.77		
35	NT2RM4001281	4	4	10.54	10.15	9.05	10		
	NT2RM4001286	345.27	345.27	526.77	215.2	220.51	552.53		
	NT2RM4001290	23.51	23.51	61.5	56.51	48.6	59.58		
40	NT2RM4001309	2.64	2.64	6.81	3.17	5.42	4.05		
	NT2RM4001313	2.63	2.63	8.62	3.81	6.36	7.03		
	NT2RM4001316	3.14	3.14	6.12	3.39	3.85	5.25		
45	NT2RM4001320	2.4	2.4	6.43	2.83	3.16	5.6		
	NT2RM4001321	3.98	3.98	8.62	6.17	7.03	6.24		
	NT2RM4001325	2.54	2.54	5.2	4.76	2.93	5.25		
	NT2RM4001333	8.65	8.65	18.06	8.57	10.76	9.51		
50	NT2RM4001340	4.81	4.81	12.27	6.99	7.72	11.24		
	NT2RM4001344	4.09	4.09	4.69	3.04	3.44	4.93		
	NT2RM4001347	6.49	6.49	9.8	10.53	9.72	17.71		
	NT2RM4001357	7.59	7.59	12.09	8.58	11.68	9.37		
55	NT2RM4001360	2.79	2.79	5.11	3.82	2.85	3.07		

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	NT2RM4000833	2.51	2.51	6.84	4.48	3.35	4.87		
	NT2RM4000841	4	4	15	10.57	8.84	10.1		
5	NT2RM4000846	1.66	1.66	8.83	4.74	6.83	5.09		
	NT2RM4000848	2.61	2.61	5.75	4.15	6.32	3.12		
	NT2RM4000852	3.89	3.89	9.81	8.16	8.67	8.29		
	NT2RM4000855	5.12	5.12	7.64	5.99	6.08	7.19		
10	NT2RM4000859	11.18	11.18	16.28	16.48	19.12	17.62		
	NT2RM4000868	3.06	3.06	6.47	5.23	6.91	7.05		
	NT2RM4000870	4	4	9.82	7.06	3.87	6.59		
15	NT2RM4000879	1.67	1.67	6	4.15	3.31	3.11		
	NT2RM4000882	18.99	18.99	28.36	20.33	16.5	12.55		
	NT2RM4000887	2.16	2.16	6.01	3.53	4.91	1.6		
	NT2RM4000895	2.33	2.33	5.33	3.16	5.9	3.08		
20	NT2RM4000897	5.78	5.78	6.99	8.89	9.73	7.02		
	NT2RM4000901	5.22	5.22	6.41	4.37	6.2	6.12		
	NT2RM4000950	4.04	4.04	5.09	3.57	4.06	2.91		
25	NT2RM4000965	2.89	2.89	4.54	5.98	4.04	6.18		
	NT2RM4000971	2.49	2.49	4.78	4.84	4.29	7.62		
	NT2RM4000979	5.16	5.16	12.71	10.23	10.46	7.87		
	NT2RM4000987	1.9	1.9	4.59	3.64	4.22	3.21		
30	NT2RM4000989	2.51	2.51	5.17	4.2	5.16	4.99		
	NT2RM4000991	3.1	3.1	4.83	2.87	3.53	8.73		
	NT2RM4000992	3.39	3.39	5.41	3.6	4.54	3.19		
35	NT2RM4000996	6.22	6.22	7.86	7.14	8	7.28		
	NT2RM4000997	3.53	3.53	13.96	9.81	10.96	10.89		
	NT2RM4001001	26.06	26.06	55.43	37.67	34.22	54.29		
	NT2RM4001002	5.13	5.13	11.03	11.54	11.33	19.44		
40	NT2RM4001016	1.63	1.63	2.73	4.07	5.31	3.76	*	+
	NT2RM4001025	65.77	65.77	133.97	148.39	181.87	171.5	*	+
	NT2RM4001027	2.49	2.49	3.66	1.67	1.77	4.31		
45	NT2RM4001032	2.55	2.55	6.74	4.94	5.46	3.84		
	NT2RM4001047	3.87	3.87	3.7	2.61	2.73	2.7	**	-
	NT2RM4001049	3.97	3.97	10.12	18.29	20.63	26.25	**	+
	NT2RM4001051	2.72	2.72	12.54	11.17	10.12	13.68		
50	NT2RM4001052	14.95	14.95	72.14	75.49	75.01	79.12		
	NT2RM4001053	14.96	14.96	39.3	41.36	28.95	25.87		
	NT2RM4001054	3.13	3.13	5.17	5.34	5.69	5.01		
55	NT2RM4001059	3.65	3.65	6.37	4.91	3.52	4.48		
	NT2RM4001071	4.03	4.03	7.35	6.8	7.34	6.55		

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	NT2RM4000603	4.18	4.18	6.98	6.95	7.23	7.26		
	NT2RM4000611	9.81	9.81	18.02	17.1	17.38	25.49		
5	NT2RM4000616	2.05	2.05	-6.19	2.73	4.14	4.53		
	NT2RM4000621	26.04	26.04	70.86	57.5	62.2	59.07		
	NT2RM4000648	2.78	2.78	7.83	2.66	2.67	3.88		
10	NT2RM4000649	5.53	5.53	11.03	7.34	8.4	13.12		
	NT2RM4000658	3.22	3.22	8.18	4.22	6.14	8.14		
	NT2RM4000661	57.68	57.68	140.68	141.09	135.49	177.49		
	NT2RM4000673	3.02	3.02	5.72	3.69	3.8	4.59		
15	NT2RM4000674	2.23	2.23	4.27	2.89	2.63	2.99		
	NT2RM4000689	3.81	3.81	6.47	5.34	6.36	7.74		
	NT2RM4000698	14.85	14.85	20.92	25.12	23.66	26.39	*	+
20	NT2RM4000700	2.39	2.39	5.53	2.41	6.95	5.61		
	NT2RM4000701	10.07	10.07	54.36	61.81	67.75	63.17		
	NT2RM4000712	3.5	3.5	7.9	5.97	8.03	9.55		
	NT2RM4000717	2.14	2.14	6.66	3.67	2.94	4.3		
25	NT2RM4000733	4.37	4.37	7.8	4.16	6.93	11.03		
	NT2RM4000734	2.17	2.17	5.92	2.35	5.23	4.7		
	NT2RM4000741	2.14	2.14	6.11	3.59	4.75	4.66		
30	NT2RM4000744	1.76	1.76	7.05	2.76	4.4	10.18		
	NT2RM4000749	15.53	15.53	23.13	26.26	27.8	34.67	*	+
	NT2RM4000751	2.88	2.88	6.54	6.23	6.11	5.94		
	NT2RM4000752	4.11	4.11	4.88	4.78	5.12	38.58		
35	NT2RM4000760	3.5	3.5	9.69	4.54	6.31	5.26		
	NT2RM4000761	237.9	237.9	478.3	219.65	302.54	336.34		
	NT2RM4000764	66.05	66.05	178	212.33	205.98	232.75	*	+
40	NT2RM4000768	6.11	6.11	11.21	15.56	10.14	21.17		
	NT2RM4000778	1.6	1.6	4.7	4.27	5.18	6.18		
	NT2RM4000779	4.52	4.52	8.28	6.87	7.19	7.33		
	NT2RM4000787	2.55	2.55	7.49	3.64	4.9	4.53		
45	NT2RM4000790	2.99	2.99	5.03	5.47	5.82	12.06		
	NT2RM4000795	1.99	1.99	3.67	2.36	1.2	2.51		
	NT2RM4000796	3.26	3.26	5.86	4.29	3.48	4.28		
	NT2RM4000798	1.77	1.77	5.53	3.72	3.08	3.47		
50	NT2RM4000800	4.15	4.15	8.16	8.7	9.44	9.06		
	NT2RM4000813	3.31	3.31	8.79	7.14	7.95	10.09		
	NT2RM4000820	4.89	4.89	9.14	5.39	6.27	5.44		
55	NT2RM4000827	7.1	7.1	18.55	16.3	15.8	17.88		
	NT2RM4000830	3.27	3.27	7.35	5.28	7.8	8.38		

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	NT2RM4000349	3.68	3.68	9.99	11.87	10.88	13.8	*	+
	NT2RM4000354	1.65	1.65	3.13	4.2	4.31	3.1		
5	NT2RM4000356	1.5	1.5	-3.11	2.5	4.07	1.64		
	NT2RM4000366	15.75	15.75	44.48	38.81	44.07	58.06		
	NT2RM4000368	3.04	3.04	5.9	4.36	5.48	3.48		
	NT2RM4000373	6.49	6.49	12.29	12.72	15.96	16.47	*	+
10	NT2RM4000386	4.92	4.92	4.71	3.81	4.57	4.6		
	NT2RM4000395	2.7	2.7	4.69	6.36	6.51	5.68	*	+
	NT2RM4000414	1	1	2.76	2.38	2.9	2.19		
15	NT2RM4000417	1.66	1.66	2.83	3.9	3.95	3.25	*	+
	NT2RM4000421	2.99	2.99	5.17	4.96	5.47	4.13		
	NT2RM4000425	10.56	10.56	26.8	26.49	31.48	45.28		
	NT2RM4000433	2.78	2.78	5.39	1.67	2.21	1.79		
20	NT2RM4000436	3.8	3.8	9.47	11.84	16.75	16.38	*	+
	NT2RM4000444	4.51	4.51	12.97	7.29	8.54	7.38		
	NT2RM4000457	3.35	3.35	8.69	13.35	12.38	13.25	*	+
25	NT2RM4000471	1.73	1.73	4.01	4.17	4.87	2.49		
	NT2RM4000472	2.2	2.2	7.62	6.64	7.61	5.39		
	NT2RM4000486	2.98	2.98	5.92	6.85	7.54	6		
	NT2RM4000490	3.85	3.85	6.41	7.16	5.1	5.86		
30	NT2RM4000496	3.68	3.68	3.86	2.16	2.36	2.04	**	-
	NT2RM4000505	26.85	26.85	60.33	68.9	80.59	70.67	*	+
	NT2RM4000511	22.8	22.8	45.35	64.6	89.95	75.97	*	+
35	NT2RM4000514	2.61	2.61	6.75	10.47	7.53	9.25	*	+
	NT2RM4000515	3.75	3.75	8.81	7.27	7.94	4.66		
	NT2RM4000517	34.51	34.51	74.2	76.45	53.07	79.47		
	NT2RM4000520	2.24	2.24	3.08	3.21	3.3	5.49		
40	NT2RM4000531	2.76	2.76	5.71	4.41	5.5	4.22		
	NT2RM4000532	3.03	3.03	5.56	2.72	3.54	2.39		
	NT2RM4000533	3.55	3.55	5.43	4.73	6.98	3.23		
45	NT2RM4000534	5.17	5.17	2.92	2.73	4.62	1.94		
	NT2RM4000563	5.21	5.21	16.69	17.05	16.8	19.36		
	NT2RM4000566	1.79	1.79	4.6	7.14	5.27	7.91	*	+
	NT2RM4000568	10.48	10.48	19.4	28.18	22.91	33.06	*	+
50	NT2RM4000585	1.97	1.97	2.52	0.82	1.99	1.26		
	NT2RM4000587	3.88	3.88	6.15	6.66	5.75	5.15		
	NT2RM4000590	3.73	3.73	4.12	1.64	2.48	2.3	**	-
55	NT2RM4000593	4.46	4.46	7.83	10.21	12.36	7.45		
	NT2RM4000595	3.94	3.94	3.91	2.27	4.36	2.45		

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	NT2RM4000102	32.35	32.35	42.47	27.24	19.34	70.54		
	NT2RM4000104	2.78	2.78	7.13	3.41	5.12	4.3		
5	NT2RM4000115	2.87	2.87	6.1	3.91	5.86	4.08		
	NT2RM4000129	2.17	2.17	4.75	2.62	3.48	2.18		
	NT2RM4000139	3.17	3.17	3.31	3.58	4.82	6.49		
	NT2RM4000149	2.74	2.74	1.49	2.41	2.55	7.32		
10	NT2RM4000155	2.73	2.73	5.5	2.13	4.51	3.71		
	NT2RM4000156	5.94	5.94	16.74	20.45	21	21.96	*	+
	NT2RM4000167	1.36	1.36	2.58	2.61	4.34	1.75		
15	NT2RM4000169	9.95	9.95	36.53	29.12	24.89	23.57		
	NT2RM4000191	4.29	4.29	7.56	5.49	5.57	5.66		
	NT2RM4000197	2.73	2.73	4.78	1.83	3.86	2.2		
	NT2RM4000198	3.38	3.38	7.42	5.26	5.45	4.21		
20	NT2RM4000199	2	2	3.51	2.8	4	3.76		
	NT2RM4000200	0.67	0.67	3.19	2.25	1.84	1.2		
	NT2RM4000202	1	1	3.24	2.11	2.42	1.84		
25	NT2RM4000210	1.46	1.46	3.72	2.41	3.08	2.21		
	NT2RM4000215	2.54	2.54	5.43	3.3	4.09	3.06		
	NT2RM4000220	6.42	6.42	10.52	8.68	11.08	15.14		
	NT2RM4000229	3.26	3.26	6.62	1.46	2.34	1.76		
30	NT2RM4000231	6.37	6.37	7.06	6.13	7.85	6.24		
	NT2RM4000233	4.83	4.83	17.3	11.9	14.34	13.4		
	NT2RM4000244	2.35	2.35	5.22	3.86	4.14	5.17		
35	NT2RM4000251	3.85	3.85	10.97	4.82	6.52	4.59		
	NT2RM4000255	2.28	2.28	4.7	3.79	4.89	1.42		
	NT2RM4000265	2.23	2.23	5.69	4.29	8.21	1.99		
	NT2RM4000283	18.14	18.14	26.21	37.17	39.88	44.79	**	+
40	NT2RM4000284	13.85	13.85	33.72	31.96	42.88	40.67		
	NT2RM4000290	6.31	6.31	7.76	3.77	4.92	4	*	-
	NT2RM4000295	2.36	2.36	2.16	2.32	2.13	0.88		
	NT2RM4000306	3.79	3.79	7.76	7.1	6.14	5.02		
45	NT2RM4000307	5.04	5.04	9.13	9.95	9.99	11.72	*	+
	NT2RM4000309	2.48	2.48	5.34	3.92	6.52	5.52		
	NT2RM4000313	3.92	3.92	9.61	5.75	7.77	8.52		
50	NT2RM4000318	3.38	3.38	6.87	4.35	6.36	3.28		
	NT2RM4000324	4.93	4.93	5.93	2.79	4.98	2.12		
	NT2RM4000326	5.32	5.32	4.61	2.59	2.45	2.01	**	-
	NT2RM4000327	4.97	4.97	10.95	7.94	10.32	7.71		
55	NT2RM4000344	5.46	5.46	16.67	11.16	10.17	19.18		

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	NT2RM2002647	7.68	7.68	17.56	19.5	17.74	23.44		
	NT2RM2002652	5.11	5.11	4.71	4.3	6.43	3.73		
5	NT2RM2002692	4.59	4.59	4.48	2.5	2.86	2.02	**	-
	NT2RM2002721	30.26	30.26	46.01	62.71	84.18	61.02	*	+
	NT2RM2002748	18.37	18.37	43.62	87.35	119.27	102.35	**	+
	NT2RM2002764	2.28	2.28	5.3	5.05	5.95	4.07		
10	NT2RM2002772	3.15	3.15	9.32	9.66	7.81	5.44		
	NT2RM2002811	5.79	5.79	12.3	14.01	9.18	10.45		
	NT2RM2002818	2.03	2.03	7.94	5.86	5.42	7.13		
15	NT2RM2002879	4.21	4.21	7.17	8.39	7.87	9.11	*	+
	NT2RM2002979	11.79	11.79	19.66	24.49	23.23	21.79	*	+
	NT2RM2002981	4.42	4.42	3.78	3.58	4.95	2.63		
	NT2RM2002995	5.13	5.13	3.29	3.42	3.74	3.5		
20	NT2RM2003031	1.37	1.37	2.63	3	2.58	1.9		
	NT2RM2003042	4.1	4.1	10.77	10.59	6.02	6.07		
	NT2RM2003044	1.88	1.88	4.11	2.13	5.11	1.2		
25	NT2RM2003090	4.4	4.4	7.64	9.36	7.91	10.68	*	+
	NT2RM2003095	11.98	11.98	25.25	15.63	16.43	19.04		
	NT2RM2003116	11.16	11.16	16.09	17.96	21.43	22.08	*	+
	NT2RM2003222	3.98	3.98	3.63	2.67	3.64	2.35		
30	NT2RM2003224	11.29	11.29	15.33	24.29	29.77	20.76	*	+
	NT2RM2003250	14.18	14.18	86.06	85.79	96.6	94.15		
	NT2RM2003258	4.59	4.59	6.32	6.54	5.11	5.69		
35	NT2RM2003262	5.07	5.07	7.33	5.06	7.76	5.72		
	NT2RM4000023	2.15	2.15	7.02	3.57	4.91	4.29		
	NT2RM4000024	2.28	2.28	6.78	3.17	4.98	4.33		
	NT2RM4000027	4.74	4.74	7.77	4.85	6.94	11.32		
40	NT2RM4000030	2.95	2.95	5.73	3.16	4.9	2.64		
	NT2RM4000033	2.51	2.51	4.77	2.36	3.89	4.61		
	NT2RM4000034	1.93	1.93	5.35	3.74	4.84	6.09		
45	NT2RM4000046	1.37	1.37	3.79	1.57	3.5	2.32		
	NT2RM4000052	1.82	1.82	3.55	1.72	2.96	1.98		
	NT2RM4000054	10.43	10.43	13.85	12.07	12.83	24.7		
	NT2RM4000061	1.65	1.65	4.17	1.66	4.54	0.83		
50	NT2RM4000074	15.83	15.83	43.57	27.9	34.24	30.79		
	NT2RM4000085	5.35	5.35	10.1	8.41	10.19	10.17		
	NT2RM4000086	3.06	3.06	4.5	3.84	5.25	3.71		
55	NT2RM4000100	6.62	6.62	15.05	12.74	15.6	14.84		
	NT2RM4000101	3.77	3.77	9.11	7.17	7.71	8.78		

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	NT2RM2002049	7.4	7.4	12.12	9.76	10.42	13.22		
	NT2RM2002055	2.8	2.8	8.01	1.91	4.03	2.52		
5	NT2RM2002072	9.26	9.26	12.88	12.28	19.12	12.82		
	NT2RM2002088	4.82	4.82	13.85	11.35	11.7	15.28		
	NT2RM2002091	4.98	4.98	8.44	5.92	3.82	4.61		
	NT2RM2002100	3.26	3.26	6.05	4.82	4.24	3.19		
10	NT2RM2002109	1.31	1.31	3.57	2.57	4.88	4.92		
	NT2RM2002126	21.41	21.41	32.24	35.28	22.31	31.52		
	NT2RM2002128	3.7	3.7	5.17	2.74	3.86	2.41		
15	NT2RM2002129	6.43	6.43	11.48	8.53	13.03	10.66		
	NT2RM2002142	5.72	5.72	9.74	5.26	8.91	6.4		
	NT2RM2002144	3.27	3.27	3.76	1.85	1.73	1.62	**	-
	NT2RM2002145	2.63	2.63	8.69	6.1	5.18	5.98		
20	NT2RM2002153	2.61	2.61	6.37	6.31	7.62	5.75		
	NT2RM2002163	0.97	0.97	3.41	1.87	3.4	0.64		
	NT2RM2002170	3.28	3.28	7.03	6.62	7.5	7.65		
25	NT2RM2002178	3.99	3.99	3.67	3.5	5.48	2.9		
	NT2RM2002179	7.82	7.82	8.69	6.17	8.15	6.02		
	NT2RM2002270	4.51	4.51	4.56	2.28	1.76	1.67	**	-
	NT2RM2002326	2.47	2.47	3.86	2.13	3.69	2.34		
30	NT2RM2002337	1.88	1.88	3.97	5.4	4.22	4.79	*	+
	NT2RM2002339	2.83	2.83	6.29	5.26	5.22	3.85		
	NT2RM2002345	5.16	5.16	6.03	4.04	4.2	4.21	*	-
35	NT2RM2002368	2.43	2.43	5.86	6.05	7.01	4.96		
	NT2RM2002381	2.23	2.23	5.16	3.47	3.65	2.8		
	NT2RM2002424	4.64	4.64	7.1	6.69	8.5	6.3		
	NT2RM2002450	4.17	4.17	3.87	2.29	2.39	1.87	**	-
40	NT2RM2002482	3.93	3.93	4.65	2.66	3.2	3.79		
	NT2RM2002492	9.39	9.39	24.31	29.13	24.65	29.29		
	NT2RM2002575	3.26	3.26	5.23	5.99	6.03	5.07		
45	NT2RM2002580	4.23	4.23	4.68	4.82	7.79	7.42		
	NT2RM2002592	7.7	7.7	12.59	13.07	15.28	14.69	*	+
	NT2RM2002608	27.33	27.33	45.49	57.07	65.96	48.3	*	+
	NT2RM2002615	6.01	6.01	9.38	13.15	20.32	14.42	*	+
50	NT2RM2002622	14.35	14.35	16.22	18.38	24.99	13.44		
	NT2RM2002630	4.86	4.86	6.63	8.05	7.37	6.7		
	NT2RM2002634	1.72	1.72	4.66	4.71	4.6	3.94		
55	NT2RM2002645	27.02	27.02	68.46	30.66	14.59	31.46		
	NT2RM2002646	12.09	12.09	25.03	29.45	22.88	34.8		

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	NT2RM2001825	14.79	14.79	36.08	34.68	34.2	35.81		
	NT2RM2001832	5.93	5.93	6.1	5.19	5.48	2.93		
5	NT2RM2001839	67.48	67.48	123.46	152.63	157.11	121.35		
	NT2RM2001840	3.04	3.04	7.13	4.61	5.11	5.37		
	NT2RM2001851	3.92	3.92	7.61	3.78	4.74	6.49		
10	NT2RM2001855	8.21	8.21	11.51	10.22	12.06	15.41		
	NT2RM2001867	2.82	2.82	5.01	2.83	5.62	3.74		
	NT2RM2001869	60.8	60.8	90.58	101.19	79.67	105.32		
	NT2RM2001879	3.01	3.01	6.99	2.55	3.19	2.66		
15	NT2RM2001883	1.52	1.52	3.26	0.98	2.28	0.73		
	NT2RM2001886	1.57	1.57	4.56	2.48	3.51	2.42		
	NT2RM2001887	3.78	3.78	7.66	4.48	4.97	5.73		
20	NT2RM2001896	274.2	274.2	378.57	325.68	216.52	497.31		
	NT2RM2001902	1.92	1.92	4.28	1.31	2.92	2.14		
	NT2RM2001903	16.25	16.25	42.55	35.47	31.71	37.22		
	NT2RM2001930	2.11	2.11	6.3	2.3	5.86	5.44		
25	NT2RM2001935	4.16	4.16	5.04	3.16	4.42	5.87		
	NT2RM2001936	2.81	2.81	4.9	2.99	3.44	4		
	NT2RM2001939	3.56	3.56	3.34	1.82	3.01	3.16		
30	NT2RM2001941	1.84	1.84	4.29	2.84	2.82	2.72		
	NT2RM2001950	4.66	4.66	10	6.01	6	8.69		
	NT2RM2001952	2.67	2.67	4.78	2.49	4.55	5.37		
	NT2RM2001976	11.48	11.48	18.2	14.58	11.46	35.27		
35	NT2RM2001982	1.85	1.85	3.91	2.04	2.47	1.88		
	NT2RM2001983	4.45	4.45	8.36	4.18	6.49	7.54		
	NT2RM2001984	7.74	7.74	8.88	20.06	27.05	22.95	**	+
	NT2RM2001989	2.72	2.72	3.68	2.99	4.26	3.7		
40	NT2RM2001996	7.51	7.51	8.09	4.8	8.01	5.29		
	NT2RM2001997	3.65	3.65	7.29	3.18	5.09	6.78		
	NT2RM2001998	2.24	2.24	5.07	3.33	6.53	3.96		
45	NT2RM2001999	4.86	4.86	7.69	6.88	6.02	4.01		
	NT2RM2002003	11.33	11.33	18.17	10.15	11	14.9		
	NT2RM2002004	1.99	1.99	5.79	2.51	2.09	1.9		
	NT2RM2002009	5.35	5.35	9.03	9.85	11.04	11.09	*	+
50	NT2RM2002014	2.62	2.62	3	3.65	4.47	4.03	**	+
	NT2RM2002019	25.1	25.1	38.52	19.47	14.2	16.35		
	NT2RM2002029	12.92	12.92	19.01	10.82	4.88	14.74		
55	NT2RM2002030	4.15	4.15	5.8	13.54	9.71	21.98	*	+
	NT2RM2002034	22.05	22.05	31.76	24.83	20.89	21.04		

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	NT2RM2001685	4. 58	4. 58	5. 68	1. 29	2. 72	1. 14	**	-
	NT2RM2001688	5. 46	5. 46	4. 14	3. 11	3. 82	2. 46	*	-
5	NT2RM2001695	15. 09	15. 09	35. 18	17. 41	19. 26	34. 51		
	NT2RM2001696	2. 74	2. 74	6. 64	7. 15	6. 7	6. 8		
	NT2RM2001698	1. 44	1. 44	3	4. 06	3. 49	1. 65		
	NT2RM2001699	1. 63	1. 63	5. 03	4. 19	3. 75	5. 48		
10	NT2RM2001700	1. 65	1. 65	4. 13	2. 56	3. 37	3. 91		
	NT2RM2001704	2. 68	2. 68	5. 46	3. 89	3. 85	3. 99		
	NT2RM2001706	4. 29	4. 29	6. 77	3. 33	3. 13	3. 32		
15	NT2RM2001714	6. 48	6. 48	6. 64	5. 62	7. 33	5. 18		
	NT2RM2001716	0. 97	0. 97	3. 7	3. 03	5. 49	2. 92		
	NT2RM2001718	1. 91	1. 91	3. 47	5	3. 5	3		
	NT2RM2001723	2. 09	2. 09	5. 48	5. 1	5. 21	5. 71		
20	NT2RM2001727	3. 08	3. 08	6. 25	7. 51	7. 28	5. 7		
	NT2RM2001730	3. 52	3. 52	7. 15	5. 04	5. 43	3. 85		
	NT2RM2001738	4. 56	4. 56	6. 2	6. 71	10. 25	9. 08	*	+
25	NT2RM2001743	2. 95	2. 95	5. 81	4. 39	5. 02	4. 46		
	NT2RM2001753	5. 98	5. 98	7. 55	5. 72	6. 09	4. 54		
	NT2RM2001755	0. 89	0. 89	2. 82	2. 4	2. 83	2. 67		
	NT2RM2001760	14. 77	14. 77	33. 17	27. 49	25. 48	36. 23		
30	NT2RM2001765	1. 35	1. 35	1. 71	2. 45	3. 12	2. 03	*	+
	NT2RM2001767	12. 04	12. 04	120. 66	148. 84	168. 4	146. 29	*	+
	NT2RM2001768	2. 1	2. 1	3. 59	3. 41	4. 21	3. 05		
35	NT2RM2001771	4. 82	4. 82	5. 65	7. 15	5. 97	5. 05		
	NT2RM2001778	2. 89	2. 89	4. 09	2. 34	3. 24	1. 48		
	NT2RM2001782	5. 32	5. 32	7. 32	4. 96	7. 71	7. 57		
	NT2RM2001784	0. 84	0. 84	2. 19	2. 81	2. 5	1. 41		
40	NT2RM2001785	1. 35	1. 35	4. 11	5. 5	5. 02	2. 76		
	NT2RM2001792	6. 03	6. 03	8. 53	5. 49	5. 54	5. 76		
	NT2RM2001795	3. 97	3. 97	6. 15	7. 62	5. 96	8. 9		
45	NT2RM2001797	2. 82	2. 82	3. 78	5	5. 94	2. 71		
	NT2RM2001800	3. 46	3. 46	4. 26	5. 01	4. 03	5. 24		
	NT2RM2001803	3. 5	3. 5	6. 61	4. 46	7. 34	2. 44		
	NT2RM2001805	3. 65	3. 65	3. 21	2. 53	4. 2	1. 71		
50	NT2RM2001806	7. 34	7. 34	17. 96	15. 62	15. 23	21. 11		
	NT2RM2001813	1. 54	1. 54	2. 05	2. 54	1. 88	2. 32		
	NT2RM2001814	2. 46	2. 46	4. 71	3. 52	2. 89	4. 42		
55	NT2RM2001818	1. 21	1. 21	2. 66	0. 97	1. 48	0. 27		
	NT2RM2001823	1. 4	1. 4	3. 24	1. 87	2. 46	1. 37		

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	NT2RM2001423	5.44	5.44	9.53	8.64	11.95	11.36		
	NT2RM2001424	5.88	5.88	15.09	11.77	10.31	11.63		
5	NT2RM2001482	2.24	2.24	6.48	3.5	6.06	3.63		
	NT2RM2001499	1.4	1.4	5.81	2.84	4.3	2.17		
	NT2RM2001504	3.63	3.63	6.99	3.2	4.54	1.68		
10	NT2RM2001524	2.51	2.51	5.81	2.34	2.22	3.51		
	NT2RM2001530	2.56	2.56	4.42	2.68	4.35	3.52		
	NT2RM2001533	5.06	5.06	9.09	8.2	9.18	7.84		
	NT2RM2001540	5.77	5.77	8.36	14.57	17.99	27.1	*	+
15	NT2RM2001544	2.4	2.4	6.12	3.7	3.72	2.31		
	NT2RM2001547	6.6	6.6	15.29	8.44	7.61	8.24		
	NT2RM2001558	1.53	1.53	3.44	1.76	4.87	1.71		
20	NT2RM2001575	2.45	2.45	4.57	3.36	4.38	2.29		
	NT2RM2001582	2.99	2.99	4.98	2.2	5.16	3.06		
	NT2RM2001588	3.69	3.69	8.8	6.39	9.14	6.6		
	NT2RM2001592	2.66	2.66	6.2	3.1	5.24	4.64		
25	NT2RM2001603	4.74	4.74	8.7	10.42	12.03	11.77	*	+
	NT2RM2001605	1.74	1.74	4.52	3.08	1.51	2.39		
	NT2RM2001611	2.28	2.28	8.63	3.74	3.34	3.51		
30	NT2RM2001613	14.91	14.91	32.53	21.51	13.13	27.42		
	NT2RM2001626	2.45	2.45	3.08	2.1	4.28	2.06		
	NT2RM2001632	4.93	4.93	7.07	4.67	4.88	5.42		
	NT2RM2001633	4.45	4.45	10.39	3.74	5.15	5.43		
35	NT2RM2001635	4.33	4.33	9.54	4.3	5.81	4.7		
	NT2RM2001636	4.88	4.88	7.35	12.75	18.11	13.34	**	+
	NT2RM2001637	1.25	1.25	6.48	4.18	3.68	2.51		
	NT2RM2001639	3.98	3.98	9.32	4.67	4.33	3.29		
40	NT2RM2001641	1.63	1.63	4.69	4.84	6.02	2.71		
	NT2RM2001643	2.78	2.78	7.46	4.79	4.4	2.83		
	NT2RM2001648	12.97	12.97	18.91	20.13	17.07	25.5		
45	NT2RM2001652	6.32	6.32	5.65	4.29	8.13	4.46		
	NT2RM2001659	5.78	5.78	9.17	5.73	5.28	6.95		
	NT2RM2001660	3.44	3.44	3.86	2.08	2.29	2.63	**	-
	NT2RM2001664	1.24	1.24	6.12	4.51	4.89	4.8		
50	NT2RM2001668	3.72	3.72	8.16	7.66	5.72	7.02		
	NT2RM2001670	1.62	1.62	4.11	2.88	3.96	3.56		
	NT2RM2001671	2.67	2.67	5.57	3.9	6.46	4.85		
55	NT2RM2001675	1.94	1.94	4.28	1.97	3.73	0.64		
	NT2RM2001681	2.47	2.47	5.91	3.13	4.64	3.39		

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	NT2RM2001055	3.78	3.78	5.89	4.24	6.31	4.7		
	NT2RM2001065	6.17	6.17	15.91	8.51	12.12	14.21		
5	NT2RM2001075	39.81	39.81	188.21	154.65	156.72	168.68		
	NT2RM2001083	2.23	2.23	5.57	4.01	5.8	3.01		
	NT2RM2001100	10.38	10.38	93.67	95.8	113.9	97.52		
	NT2RM2001105	6.34	6.34	8.27	11.35	5.34	11.8		
10	NT2RM2001109	6.81	6.81	9.4	11.88	12.47	14.53	*	+
	NT2RM2001110	7.67	7.67	21.63	21.2	30.71	23.29		
	NT2RM2001126	6.1	6.1	6.53	5.32	6.44	7.27		
15	NT2RM2001131	5.52	5.52	40.22	21.93	29.37	20.14		
	NT2RM2001141	1.64	1.64	6.84	7.09	6.4	5.45		
	NT2RM2001152	1.63	1.63	3.27	4.42	5.77	3.02		
	NT2RM2001177	3.42	3.42	7.23	10.28	7.25	8.24		
20	NT2RM2001194	2.74	2.74	7.51	6.68	5.77	8.17		
	NT2RM2001195	3.7	3.7	8.8	6.37	7.13	6.89		
	NT2RM2001196	5.24	5.24	6.35	5.19	6.46	4.64		
25	NT2RM2001201	14.45	14.45	25.36	20.02	21.68	22.38		
	NT2RM2001221	4.22	4.22	8.61	11.69	13.61	16.63	*	+
	NT2RM2001238	2.87	2.87	5.65	3.91	3.88	1.96		
	NT2RM2001243	5.39	5.39	8.98	9.81	6.13	6.53		
30	NT2RM2001244	3.91	3.91	10.63	6.58	9.24	6.41		
	NT2RM2001247	14.94	14.94	121.59	110.47	140.27	118.79		
	NT2RM2001256	3.84	3.84	5.23	3.15	3.26	2.96		
35	NT2RM2001269	4.4	4.4	5.98	4.8	5.63	4.74		
	NT2RM2001278	5.28	5.28	7.37	8.45	8.56	5.35		
	NT2RM2001291	3.05	3.05	5.18	3.24	4.62	2.9		
	NT2RM2001294	12.47	12.47	24.39	20.08	15.43	17.81		
40	NT2RM2001295	2.56	2.56	8.82	4.54	4.43	4.99		
	NT2RM2001302	2.38	2.38	4.55	2.3	4.5	2.81		
	NT2RM2001306	3.51	3.51	7.62	4.1	4.46	5.14		
45	NT2RM2001312	2.34	2.34	3.72	1.92	2.84	1.68		
	NT2RM2001319	2.76	2.76	3.93	3.61	5.29	4.11		
	NT2RM2001324	3.73	3.73	8.29	5.48	4.9	5.71		
	NT2RM2001345	8.53	8.53	10.01	6.83	11.12	14.14		
50	NT2RM2001360	4.02	4.02	6.36	5.67	5.9	5.46		
	NT2RM2001370	5.75	5.75	14.53	8.56	9.86	11.69		
	NT2RM2001391	1.79	1.79	6.07	1.85	5.04	1.65		
55	NT2RM2001393	4.49	4.49	6.39	5.12	7.91	7.14		
	NT2RM2001420	2.94	2.94	4.61	2.61	3.62	3.14		

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	NT2RM2000660	11.45	11.45	18.34	17.03	7.1	20.16		
	NT2RM2000669	3.6	3.6	6.51	5.28	4.28	6.69		
5	NT2RM2000689	31.07	31.07	-59.7	37.03	16.51	70.9		
	NT2RM2000691	2.09	2.09	5.73	4.83	7.13	4.27		
	NT2RM2000714	3.41	3.41	10.97	11.46	14.54	11.3		
	NT2RM2000718	4.08	4.08	7.15	2.88	5.42	4.33		
10	NT2RM2000732	5.38	5.38	14.81	9.49	14.18	8.25		
	NT2RM2000735	3.72	3.72	6.16	4.27	6.55	6.49		
	NT2RM2000740	2.26	2.26	6.2	4.27	3.01	3.71		
15	NT2RM2000743	2.26	2.26	7.89	5.65	3.24	3.89		
	NT2RM2000772	6.43	6.43	8.48	5.24	6.72	9.47		
	NT2RM2000773	8.17	8.17	19.56	19.18	17.96	18.29		
	NT2RM2000776	13.96	13.96	17.16	24.24	9.95	26.76		
20	NT2RM2000784	6.64	6.64	8.8	8.74	9.02	10.5		
	NT2RM2000795	4.35	4.35	13.56	7.44	8.66	10.45		
	NT2RM2000796	2.27	2.27	4.64	1.71	2.31	1.38		
25	NT2RM2000798	25.81	25.81	160.08	158.19	136.83	188.99		
	NT2RM2000801	45.09	45.09	161.29	160.44	152.13	189.56		
	NT2RM2000821	7.53	7.53	12.33	7.37	7.77	11.87		
	NT2RM2000829	5.76	5.76	13.01	8.05	10.13	11.75		
30	NT2RM2000837	3.29	3.29	7.28	4.27	6.08	4.18		
	NT2RM2000924	9.96	9.96	36.74	43.24	57.8	35.84		
	NT2RM2000930	10.64	10.64	18.29	24.45	27.78	28.34	**	+
35	NT2RM2000937	4.35	4.35	8.62	5.08	6.66	6.56		
	NT2RM2000939	1.12	1.12	2.37	2.67	1.84	1.82		
	NT2RM2000942	124.8	124.8	253.61	161.4	118.61	210.11		
	NT2RM2000951	1.01	1.01	2.6	2.13	3.12	1.6		
40	NT2RM2000952	2.53	2.53	5.31	5.98	6.41	6.86	*	+
	NT2RM2000966	19.69	19.69	111.88	95.61	137.32	135.9		
	NT2RM2000973	23.45	23.45	16.81	39.12	39.51	33.8	**	+
45	NT2RM2000983	10.07	10.07	18.59	30.68	39.13	27.52	*	+
	NT2RM2000984	6.48	6.48	7.71	4.88	5.64	5.26	*	-
	NT2RM2000994	8.27	8.27	16.47	13.2	8.36	21.81		
	NT2RM2001004	6.01	6.01	48.58	47.51	54.15	46.8		
50	NT2RM2001022	101.09	101.09	350.92	239.63	304.37	490.04		
	NT2RM2001035	10.75	10.75	24.98	24.17	24.54	34.51		
	NT2RM2001038	5.77	5.77	9.86	10.48	11.75	7.9		
55	NT2RM2001043	4.45	4.45	10.02	5.1	7.41	6.74		
	NT2RM2001050	2.71	2.71	6.89	4.72	5.83	4.03		

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	NT2RM2000402	12.38	11	15.78	25.15	18.31	22.51	*	+
	NT2RM2000405	1.33	1.25	2.2	1.52	3.08	3.16		
5	NT2RM2000407	0.76	1.78	2.49	1.89	2.72	2.89		
	NT2RM2000410	0.79	1.94	2.23	1.98	2.84	2.09		
	NT2RM2000420	3.09	2.52	4.43	4.24	4.5	3.26		
	NT2RM2000422	3.22	2.44	5.81	3.61	6.17	2.87		
10	NT2RM2000423	1.91	1.96	5.69	3.89	7.64	4.18		
	NT2RM2000452	3.46	3.18	4.31	7.35	8.65	9.57	**	+
	NT2RM2000469	3.28	3.28	4.44	1.87	2.33	2.46	*	-
15	NT2RM2000490	6.03	6.03	9.18	5.55	6.16	6.9		
	NT2RM2000497	3.29	3.29	4.59	3.15	5.48	2.43		
	NT2RM2000502	4.69	4.69	10.24	5.87	7.08	7.02		
	NT2RM2000504	7.37	7.37	12.93	10.83	4.49	11.2		
20	NT2RM2000514	2.75	2.75	6.23	3.11	3.32	3.8		
	NT2RM2000522	1.9	1.9	3.27	1.94	1.18	1.13		
	NT2RM2000540	6.02	6.02	9.53	9.12	8.96	8.14		
25	NT2RM2000556	2.09	2.09	2.8	1.24	2.33	0.93		
	NT2RM2000565	3.35	3.35	6.02	3.27	4.14	3.72		
	NT2RM2000566	6.59	6.59	15.8	9.09	9.21	9.57		
	NT2RM2000567	2.16	2.16	5.64	2	5.67	3.82		
30	NT2RM2000569	4.69	4.69	7.93	5.77	8.18	4.7		
	NT2RM2000577	11.08	11.08	15.39	11.79	14.95	14.48		
	NT2RM2000581	4.64	4.64	6.49	5.98	7.97	6.85		
35	NT2RM2000582	5.23	5.23	10.34	8.34	9.14	7.19		
	NT2RM2000588	21.84	21.84	65.91	40.15	44.01	45.21		
	NT2RM2000589	3.98	3.98	11.35	7.96	7.6	8.64		
	NT2RM2000594	1.87	1.87	4.38	1.62	2.71	1.92		
40	NT2RM2000599	6.34	6.34	16.12	17.82	14.5	15.44		
	NT2RM2000609	4.61	4.61	6.77	3.76	5.81	5.48		
	NT2RM2000612	3.52	3.52	6.4	5.93	7.47	4.55		
45	NT2RM2000622	16.6	16.6	56.24	53.07	75.02	55.48		
	NT2RM2000623	2.66	2.66	7.1	7.92	6.03	5.58		
	NT2RM2000624	4.18	4.18	10.6	7.33	14.39	7.56		
	NT2RM2000632	2.8	2.8	6.73	4.1	6.4	4.73		
50	NT2RM2000635	3.42	3.42	8.09	5.41	6.29	5.31		
	NT2RM2000636	2.61	2.61	6.28	3.99	4.39	3.72		
	NT2RM2000639	3.73	3.73	8.26	5.42	7.79	5.99		
55	NT2RM2000649	6.03	6.03	9.69	9.4	9.17	8.05		
	NT2RM2000658	6.49	6.49	13.18	15.17	14.66	15.83	*	+

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	NT2RM1001136	0.88	1.41	2.71	2.31	3.87	1.59		
	NT2RM1001139	3.9	3.7	5.38	5.33	11.18	6.77		
5	NT2RM2000003	2.45	3.33	2.4	4.5	6.29	4.88	*	+
	NT2RM2000006	2.34	2.95	7.25	5.12	7.11	6.24		
	NT2RM2000010	12.79	13.03	22.58	20.2	17.11	21.83		
	NT2RM2000013	8.1	9.44	50.36	68.75	95.32	74.36	*	+
10	NT2RM2000030	4.8	2.21	23.41	26.33	32.15	28.69		
	NT2RM2000032	2.76	2.92	8.53	10.01	12.19	10.67	*	+
	NT2RM2000039	3.94	4.67	4.75	6.42	5.78	4.99		
15	NT2RM2000042	3.5	4.9	11.69	17.71	17.4	15.02	*	+
	NT2RM2000092	1	2.38	1.98	1.29	4.69	2.25		
	NT2RM2000093	8.37	6.63	11.41	9.02	12.23	10.18		
	NT2RM2000101	9.2	9.94	40	61.09	76.38	69.62	*	+
20	NT2RM2000104	6.82	8.02	46.75	51.34	68.83	43.48		
	NT2RM2000124	1.54	2.23	6.33	7.73	8.84	8.47	*	+
	NT2RM2000155	5.08	3.77	5.8	9.45	11.58	12.51	**	+
25	NT2RM2000191	3.33	5.68	28.62	26.54	34.38	31.6		
	NT2RM2000192	1.03	1.29	2.45	6.3	4.75	3.83	*	+
	NT2RM2000239	1.92	2.79	3.09	2.85	5.02	3.1		
	NT2RM2000240	32.78	29.59	74.35	61.15	60.54	61.71		
30	NT2RM2000241	4.49	5.9	6.35	8.24	11.72	6.78		
	NT2RM2000250	1.29	1.54	4.16	2.09	5.05	2.54		
	NT2RM2000259	3.06	3.42	3.59	6.38	8.44	6.74	**	+
35	NT2RM2000260	2.53	2.05	3.12	4.23	4.07	5.79	*	+
	NT2RM2000265	0.91	1.55	0.99	1.43	2.4	1.09		
	NT2RM2000287	4.7	4.23	10.82	10.69	11.54	14.73		
	NT2RM2000306	12.24	9.36	10.48	23.63	14	20.79	*	+
40	NT2RM2000312	19.4	17.81	25.01	38.39	31.27	24.8		
	NT2RM2000322	1.93	1.82	4.48	3.79	7.05	3.32		
	NT2RM2000343	7.74	8.38	41.34	63.81	79.6	71.12	*	+
45	NT2RM2000359	3.67	2.86	4.95	4.93	9.55	4.72		
	NT2RM2000362	20.09	18.2	62.29	94.88	111.25	95.66	*	+
	NT2RM2000363	1.08	1.89	2.97	4.2	4.32	3.33	*	+
	NT2RM2000368	2.84	2.4	4.74	6.15	5.98	5.29	*	+
50	NT2RM2000371	76.64	65.68	119.32	135.82	125	44.64		
	NT2RM2000374	1.68	1.92	5.75	3.34	4.8	3.58		
	NT2RM2000387	8.98	9.83	11.92	20.02	25.18	17.11	*	+
55	NT2RM2000393	1.7	1.63	3.75	3.31	7.65	3.28		
	NT2RM2000395	1.07	1.51	1.98	1.72	4.34	2.23		

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	NT2RM1000922	4.7	6.11	8.09	9.03	5.21	6.36		
	NT2RM1000924	0.89	3.03	3.04	3.08	2.89	3		
5	NT2RM1000927	1.35	1.78	-2.85	3.07	4.72	3.46		
	NT2RM1000951	7.95	11.33	26.73	32.33	34.46	31.18	*	+
	NT2RM1000956	7.91	6.36	13.35	23.61	27.46	21.91	**	+
	NT2RM1000960	12.48	10.27	29.06	34.95	37.47	38.96	*	+
10	NT2RM1000961	3.28	3.61	7.45	9.44	13.18	8.11		
	NT2RM1000962	4.14	3.5	8.18	7.59	10.15	9.86		
	NT2RM1000973	16.71	15.79	29.32	31.15	11.56	27.73		
15	NT2RM1000978	0.57	1.46	1.58	0.95	2.64	0.44		
	NT2RM1000982	2.34	2.29	3.52	3.57	4.94	4.54	*	+
	NT2RM1000991	1.61	1.78	4.25	3.88	5.56	5.23		
	NT2RM1000994	6.36	6.16	12.57	16.52	16.64	14.53	*	+
20	NT2RM1001002	5.11	6.69	15.34	21.78	22.69	22.28	*	+
	NT2RM1001003	5.42	5.15	11.98	16.24	9.06	8.46		
	NT2RM1001008	1.4	2.22	2.48	1.83	4.34	4.33		
25	NT2RM1001011	6.29	5.43	7.86	14.4	10.46	14.72	*	+
	NT2RM1001013	2.9	2.75	4.75	8.29	7.96	5.81	*	+
	NT2RM1001017	1	1.82	3.44	3.28	4.86	3.92		
	NT2RM1001018	65.15	74.45	146.86	134.65	125.46	113.93		
30	NT2RM1001026	1.37	2.64	3.17	2.99	4.61	3.31		
	NT2RM1001028	0.98	1.73	2.91	1.74	1.89	0.76		
	NT2RM1001043	4.47	3.64	8.42	11.43	12.7	8.01		
35	NT2RM1001044	2.23	3.17	4.92	5.03	5.51	3.93		
	NT2RM1001059	1.47	3.72	4.12	4.05	6.11	3.02		
	NT2RM1001063	4.11	3.29	6.1	4.22	5.64	5.6		
	NT2RM1001066	0.86	1.85	2.44	2.23	3.99	2.85		
40	NT2RM1001072	1.8	2.8	4.33	1.94	3.74	1.52		
	NT2RM1001074	1.66	2.38	5.18	5.18	4.19	2.67		
	NT2RM1001076	1.39	2.2	4.94	3.43	4.42	1.72		
45	NT2RM1001082	1.79	2.6	5.23	5.31	5.92	4.57		
	NT2RM1001085	1.25	1.65	2.81	1.16	3.27	1.17		
	NT2RM1001092	3.82	4.2	5.57	9.34	7.94	9.82	**	+
	NT2RM1001102	1.7	2.3	4.4	2.49	5.94	4.64		
50	NT2RM1001103	4.37	3.88	7.18	6.25	10.28	8.08		
	NT2RM1001105	1.77	2.02	4.63	2.49	5.11	3.51		
	NT2RM1001112	2.68	2.66	3.69	3.85	4.75	2.43		
55	NT2RM1001115	1.44	1.57	4.72	3	6.46	3.73		
	NT2RM1001122	2.84	3.35	7.3	9.43	9.75	9.54	*	+

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	NT2RM1000747	23.34	23.92	46.23	44.66	50.12	55.15		
	NT2RM1000752	3.83	2.36	4.62	3.95	4.88	3.46		
5	NT2RM1000767	4.14	7.27	35.27	25.27	38.02	28.81		
	NT2RM1000770	2.97	3.08	6.36	4.71	6.71	5.67		
	NT2RM1000772	0.76	0.7	1.07	1.34	1.69	0.44		
10	NT2RM1000779	13.03	12.11	42.22	53.91	45.61	66.73	*	+
	NT2RM1000780	1.16	2.9	3.74	3.09	4.32	3.01		
	NT2RM1000781	1.07	0.98	1.71	2.58	4.4	1.93		
	NT2RM1000789	5.28	5.15	29.74	29.63	46.72	36.53		
15	NT2RM1000800	2.87	2.63	6.37	5.66	9.57	6.96		
	NT2RM1000802	2.44	2.99	7.5	4.34	5.47	4.82		
	NT2RM1000811	1.78	1.6	2.13	2.26	4.96	2.76		
20	NT2RM1000826	6.06	6.36	13.34	14.42	20.73	20.98	*	+
	NT2RM1000829	3.91	2.87	6.39	6.73	8.48	8.41	*	+
	NT2RM1000831	81.54	64.45	185.14	182.43	179.79	197.27		
	NT2RM1000833	14.58	13.33	42.25	76.74	73.25	67.48	**	+
25	NT2RM1000834	4.06	3.09	6.2	8.49	9.42	10.49	**	+
	NT2RM1000841	12.34	10.01	21.15	34.98	36.63	30.81	**	+
	NT2RM1000848	4.79	4.42	6.44	9.36	12.74	10.45	**	+
30	NT2RM1000850	2.66	3.42	13.41	8.55	11.79	9.74		
	NT2RM1000852	1.34	1.94	3.23	3.01	5.76	2.61		
	NT2RM1000853	1.19	2.85	2.15	3.11	3.26	3.23		
	NT2RM1000855	29.27	24.82	45.19	52.48	45.32	58.45		
35	NT2RM1000857	4.63	5	10.67	8.76	11.3	10.76		
	NT2RM1000858	7.3	7.6	15.86	9.09	11.56	10.93		
	NT2RM1000867	19.42	15.85	28.1	32.52	35.03	24.06		
40	NT2RM1000874	3.15	2.65	7.03	5.17	9.62	5.31		
	NT2RM1000882	2.36	1.37	3.71	5.39	9.31	5.45	*	+
	NT2RM1000883	5.21	3.34	7.42	5.18	11.3	7.9		
	NT2RM1000885	3.86	4.43	9.4	7.59	8.15	9.8		
45	NT2RM1000893	3.15	3.41	8.14	7.73	6.17	8.39		
	NT2RM1000894	3.29	4.4	6.18	8.14	6.11	6.61		
	NT2RM1000898	3.72	7.33	10.02	13.4	17.51	12.41	*	+
	NT2RM1000899	1.02	2.22	3.07	3.68	7.49	4.69		
50	NT2RM1000905	11.92	17.41	30.36	37.19	45.16	37.3	*	+
	NT2RM1000910	7.5	8.78	20.16	36.37	36.98	37.5	**	+
	NT2RM1000914	6.46	7.69	19.74	14.28	17.33	17.77		
55	NT2RM1000919	6.1	3.92	9.91	14.61	17.49	15.37	**	+
	NT2RM1000921	0.72	1.9	3.69	2.79	4.27	3.32		

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	NT2RM1000527	7.97	8.92	37.68	55.15	60.19	46.68	*	+
	NT2RM1000539	3.45	3.59	12.93	15.52	17.01	18.1	*	+
5	NT2RM1000542	0.85	1.05	2.99	1.17	2.35	1.02		
	NT2RM1000553	3.7	2.42	22.32	42.83	42.96	34.5	*	+
	NT2RM1000555	11.3	11.6	23.97	34.11	29.67	22.76		
	NT2RM1000558	2.09	5.34	9.74	9.56	16.24	14.29		
10	NT2RM1000563	1.47	2.42	3.36	4.07	5.58	3.95	*	+
	NT2RM1000566	0.88	1.57	3.5	3.62	6.01	2.79		
	NT2RM1000570	96.92	77.32	137.63	167.35	105.47	174.1		
15	NT2RM1000571	13.21	11.87	22.51	43.87	40.18	28.45	*	+
	NT2RM1000574	0.84	2.15	2.55	2.15	3.07	1.67		
	NT2RM1000580	1.37	2.18	4.07	5.15	7.98	2.96		
	NT2RM1000620	2.61	2.95	8.2	8.35	9.58	7.26		
20	NT2RM1000623	1.25	1.2	2.38	1.75	2.81	0.62		
	NT2RM1000630	0.79	2.28	2.39	1.68	3.51	1.67		
	NT2RM1000633	30.97	39.36	36.34	54.43	44.6	43.59	*	+
25	NT2RM1000634	1.91	4.16	8.12	2.56	7.05	5.57		
	NT2RM1000642	3.85	5.37	8.13	8.21	8.56	8.52		
	NT2RM1000647	41.3	39.09	62.11	57.72	68.29	62.69		
	NT2RM1000648	2.49	2.65	4.61	6.14	5.63	4.51		
30	NT2RM1000650	2.46	3.05	7.6	5.4	6.07	6		
	NT2RM1000661	4.48	5.7	15.82	15.48	13.45	13.18		
	NT2RM1000666	1	1.77	1.99	1.37	2.8	0.71		
35	NT2RM1000669	3.51	2.76	4.67	3.63	5.42	3.28		
	NT2RM1000672	2.23	3.95	7.81	3.98	8.47	7.22		
	NT2RM1000681	99.53	86.09	118.7	105.41	90.59	124.14		
	NT2RM1000691	2.02	2.61	5.74	3.61	7.69	3.76		
40	NT2RM1000698	1.11	1.43	4	6.42	6.29	4.11	*	+
	NT2RM1000699	1.85	2.86	3.17	3.67	4.35	4.15	*	+
	NT2RM1000702	3.71	4.64	9.47	9.31	9.72	11.4		
45	NT2RM1000703	11.56	12.36	25.24	26.72	20.42	21.06		
	NT2RM1000704	24.48	23	32.91	46.54	24.13	40.82		
	NT2RM1000725	60.92	59.45	88.28	94.89	82.36	105.67		
	NT2RM1000726	1.85	2.02	5.75	1.97	4.8	4		
50	NT2RM1000731	1.11	2.24	4.98	2.45	3.47	3.43		
	NT2RM1000741	1.38	1.87	3.16	2.69	4.15	2.9		
	NT2RM1000742	2.61	4.6	7.41	9.55	10.94	9.84	*	+
55	NT2RM1000744	2.1	3.61	7.14	4.05	5.24	5.05		
	NT2RM1000746	2.25	2.47	2.95	2.22	4.01	3.89		

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	NT2RM1000224	8.8	8.63	15.2	15.51	21.29	17.61		
	NT2RM1000236	30.38	24.19	61.14	72.86	82.44	71.5	*	+
5	NT2RM1000242	0.23	1.17	1.32	0.39	2.27	0.12		
	NT2RM1000244	1.41	1.48	3.43	6.9	3.69	6.7	*	+
	NT2RM1000252	1.75	1.5	3.4	3.4	3.06	3.18		
10	NT2RM1000256	7.88	5.89	9.46	26.12	29.45	36.8	**	+
	NT2RM1000257	1.98	3.01	5.09	4.64	6.83	6.65		
	NT2RM1000260	7.9	7.01	13.32	9.18	12.49	11.77		
	NT2RM1000269	3.87	2.87	5.12	6.63	9.78	3.87		
15	NT2RM1000271	0.71	0.8	1.87	0.46	2.47	0.51		
	NT2RM1000272	117.67	92.26	202.95	249.32	333.98	356.74	*	+
	NT2RM1000273	10.03	9.45	20.12	22.32	16.68	15.76		
20	NT2RM1000274	63.11	66.41	123.01	137.14	91.97	104.48		
	NT2RM1000280	3.95	4.18	8.18	6.71	8.72	7.93		
	NT2RM1000295	0.49	1	2.2	1.12	3.16	0.87		
	NT2RM1000300	1.51	1.87	2.78	3.63	5.75	3.09		
25	NT2RM1000304	58.38	98.72	161.87	187.58	185.55	204.78	*	+
	NT2RM1000314	1.8	2.12	3.6	3.84	4.07	4.33		
	NT2RM1000318	12.6	14.04	20.81	35.01	29.96	29.8	**	+
30	NT2RM1000335	2.76	2.57	4.34	6.29	5.41	4.09		
	NT2RM1000341	0.46	1.27	1.95	1.41	2.33	0.99		
	NT2RM1000350	3.04	3.47	5.52	7.32	5.63	6.44		
	NT2RM1000354	0.55	1.31	1.31	5.43	7.2	5.72	**	+
35	NT2RM1000355	30.24	31.5	56.85	74.62	50.25	61.33		
	NT2RM1000361	3.63	3.87	7.23	14.39	20.29	18.78	**	+
	NT2RM1000365	0.58	1.08	1.71	1.27	1.82	0.52		
40	NT2RM1000372	14.99	19.56	30.06	42.71	46.67	45.44	**	+
	NT2RM1000377	2.04	2.18	9.66	13.38	14.74	13.48	*	+
	NT2RM1000388	0.35	1.57	3.01	2.2	3.8	2.42		
	NT2RM1000394	0.45	1.31	1.87	1.43	2.72	0.69		
45	NT2RM1000399	0.53	1.57	3.25	1.98	3.2	1.81		
	NT2RM1000407	1.13	1.52	2.17	1.02	2.7	1.51		
	NT2RM1000421	0.84	0.57	2.78	1.06	1.77	1.13		
50	NT2RM1000422	20.65	23.31	54.69	87.5	82.91	79.47	*	+
	NT2RM1000430	1.22	1.57	2.01	3.2	3.67	2.95	**	+
	NT2RM1000462	1.55	2.33	7.32	5.59	7.28	8.16		
	NT2RM1000499	1.36	2.09	4.74	5	6.16	6.37	*	+
55	NT2RM1000512	12.49	13.22	19.22	10.54	14.15	19.84		
	NT2RM1000519	33.96	37.54	55.78	31.14	29.25	47.55		

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	MAMMA1003166	3.62	3.5	5.77	6.24	8.66	6.12		
	NB9N31000010	2.5	3.88	7.58	9.63	12.26	9.5	*	+
5	NB9N31000016	0.73	2.8	5.21	4.04	4.41	3.15		
	NB9N31000043	8.1	8.88	19.71	12.51	12.3	12.64		
	NB9N31000045	167.24	153.32	255.96	401.78	320.53	296.06	*	+
10	NB9N31000054	7.29	4.42	11.75	11.15	11.87	13.43		
	NB9N31000076	2.31	1.94	3.51	4.5	6.35	4.44	*	+
	NB9N31000086	2.62	2.65	6.23	3.61	9.71	7.69		
	NT2RM1000001	2.56	2.45	6.24	5.7	7.05	6.32		
15	NT2RM1000018	3.84	4.69	10.6	6.58	9.09	6.92		
	NT2RM1000032	1.12	2.64	3.88	2.28	4.92	3.21		
	NT2RM1000035	1.72	3.68	5.53	5.44	5.21	5.98		
20	NT2RM1000037	1.38	2.98	2.75	2.41	4.15	2.11		
	NT2RM1000039	3.45	5.13	5.9	6.51	7.26	8.4	*	+
	NT2RM1000042	33.96	32.7	65.25	57.46	67.15	64.39		
	NT2RM1000055	0.85	1.74	3.34	1.16	3.55	1.16		
25	NT2RM1000059	3.26	3.16	7.66	4.69	5.97	5.78		
	NT2RM1000062	1.13	1.21	1.9	3.47	4.33	2.46	*	+
	NT2RM1000065	23.8	16.41	34.06	36.15	35.1	51.38		
30	NT2RM1000066	4.13	4.31	8.98	7.23	10.95	9.81		
	NT2RM1000071	49.63	37.81	86.71	73.04	63.32	84.05		
	NT2RM1000080	1.37	2.04	3.8	5.1	5.94	4.5	*	+
	NT2RM1000086	4.04	4.65	4.08	5.01	6.23	5.58	*	+
35	NT2RM1000092	6.17	6.93	15.76	14.48	25.91	15.13		
	NT2RM1000118	0.63	1.12	1.22	0.63	1.7	0.44		
	NT2RM1000119	1.32	2.27	1.96	1.84	3.38	2.99		
40	NT2RM1000121	1.13	1.84	1.76	2.92	3.84	2.78	*	+
	NT2RM1000122	3.5	3.78	7.34	5.5	8.86	9.57		
	NT2RM1000127	0.69	1.34	1.47	2.14	3.36	3.32	*	+
	NT2RM1000131	0.71	1.7	1.47	1.36	3.02	2.53		
45	NT2RM1000132	3.2	4.88	4.83	6.86	6.46	6.31	*	+
	NT2RM1000153	1.75	1.9	3.68	2.38	4.45	4.84		
	NT2RM1000184	72.82	77.46	151.91	106.39	163.07	125.55		
50	NT2RM1000186	1.55	1.46	4.32	2.67	4.72	3.94		
	NT2RM1000187	3.11	1.96	5.16	10.09	9.1	8.78	**	+
	NT2RM1000199	1.12	1.37	2.11	2.41	3.51	2.72	*	+
	NT2RM1000213	1.32	1.75	2.38	2.66	2.71	2.22		
55	NT2RM1000215	10.95	11.07	17.21	19.51	22.84	15.14		
	NT2RM1000218	9.72	9.95	23.71	26.94	24.74	29.21		

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	MAMMA1003007	0.69	1.16	2.73	1.88	3.7	2.32		
	MAMMA1003011	1.56	1.8	3.67	3.77	5.41	3.94		
5	MAMMA1003013	3.67	5.57	39.41	47.56	59.11	54.29	*	+
	MAMMA1003015	1.16	1.8	2.21	2.54	2.9	2.19		
	MAMMA1003019	0.6	1.61	2.1	3.12	4.61	2.63		
10	MAMMA1003020	2.96	4.19	5.34	11.31	10.33	10.09	**	+
	MAMMA1003026	1.29	1.56	2.95	2.66	4.25	2.25		
	MAMMA1003031	0.61	1.71	5.64	4.13	5.85	5.89		
	MAMMA1003033	1.34	1.65	4.13	2.84	5.11	3.64		
15	MAMMA1003035	1.66	2.5	5.44	5.12	7.03	4.9		
	MAMMA1003039	0.95	0.75	3.31	2.15	4.73	2.48		
	MAMMA1003040	1.38	2.54	5.32	4.57	7.47	7.43		
20	MAMMA1003044	2.36	2.96	6.52	4.29	6.41	5.99		
	MAMMA1003047	1.82	3.67	7.61	5.74	7.05	7.13		
	MAMMA1003049	0.47	1.72	2.03	1.08	1.56	1.45		
	MAMMA1003055	1.24	1.67	4.92	3.77	5.14	3.44		
25	MAMMA1003056	0.9	0.91	1.85	1.22	2.26	1.02		
	MAMMA1003057	2.53	3.34	6.76	7.25	9.2	5.01		
	MAMMA1003066	1.65	2.06	4.73	4.1	7.08	5.07		
30	MAMMA1003075	1.11	1.71	3.16	1.85	4.37	2.32		
	MAMMA1003089	1.69	2.11	7.13	7.85	8.66	7.43		
	MAMMA1003092	1.25	1.79	3.21	2.62	4.08	1.76		
	MAMMA1003095	2.27	3.33	5.4	7.24	8.57	5.34		
35	MAMMA1003099	1.88	2.51	4.95	4.09	6.45	4.35		
	MAMMA1003102	1.33	2.04	2.88	3.2	3.27	2.39		
	MAMMA1003104	0.64	1.07	3.17	2.15	3.25	1.56		
40	MAMMA1003113	4.22	4.21	6.98	9.22	7.02	7.07		
	MAMMA1003126	12.93	14.72	20.89	19.28	12	15.63		
	MAMMA1003127	2.95	3.14	5.91	3.88	6.12	5.19		
	MAMMA1003131	2.82	3.51	4.86	3.82	5.91	5.88		
45	MAMMA1003135	3.66	4.65	7.61	2.33	4.04	2.64		
	MAMMA1003140	0.73	2.01	3.59	2.3	3.32	1.89		
	MAMMA1003146	2.08	2.24	3.89	3.17	5.09	3.3		
50	MAMMA1003150	1.18	1.8	3.01	3.37	4.29	3.45	*	+
	MAMMA1003154	0.54	1.41	2.29	2.21	3.57	2.34		
	MAMMA1003155	8.08	9.18	20.88	21.41	17.68	20.56		
	MAMMA1003157	5.94	4.82	6.07	5.18	7.05	7.89		
55	MAMMA1003163	1.74	1.69	4.23	2.55	5.46	3.08		
	MAMMA1003164	2.94	4.56	6.23	4.08	9.9	8.18		

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	MAMMA1002863	2.17	2.83	6.91	3.51	5.12	3.96		
	MAMMA1002868	2.73	3.7	6.26	6.35	9.53	10.25	*	+
5	MAMMA1002869	5.43	6.83	-26.64	22.68	30.03	29.85		
	MAMMA1002871	0.61	1.7	1.78	1.9	3.8	1.97		
	MAMMA1002875	1.9	2.59	3.99	4.48	6.35	4.06		
	MAMMA1002879	8.42	9.2	14.19	22.55	23.63	27.96	**	+
10	MAMMA1002880	1.23	2.02	2.12	1.48	5.42	2.03		
	MAMMA1002881	1.21	1.43	1.84	3.01	5.43	2.46		
	MAMMA1002885	0.96	1.59	2.71	2.6	3.26	1.59		
15	MAMMA1002886	2.63	2.52	3.9	6.01	5.37	7.05	**	+
	MAMMA1002887	1.28	1.83	2.78	2.98	5.14	4.32	*	+
	MAMMA1002890	0.79	1.7	4.05	4.39	4.8	4.01		
	MAMMA1002892	1.35	2.45	4.98	6.64	6.24	5.84	*	+
20	MAMMA1002893	4.52	3.58	5.4	7.6	8.03	8.43	**	+
	MAMMA1002895	1.43	1.31	3.28	1.81	3.89	1.64		
	MAMMA1002898	0.53	1.67	4.15	2.69	4.72	1.42		
25	MAMMA1002905	1.32	1.58	2.51	4.1	5.01	3.87	**	+
	MAMMA1002906	15.12	10.76	15.42	19.47	13.76	15.58		
	MAMMA1002908	0.99	1.24	4.28	3.53	4.24	4.07		
	MAMMA1002909	1.92	2.64	5.67	6.82	8.18	6.57	*	+
30	MAMMA1002918	2.75	2.69	5.42	5.27	7.26	6.58		
	MAMMA1002925	92.88	85.77	163.7	127.31	122.97	178.98		
	MAMMA1002926	6.08	6.31	16.25	16.64	19.48	19.9		
35	MAMMA1002930	1.21	1.59	5.67	4.88	8.91	4.21		
	MAMMA1002937	4.91	3.87	30.71	40.45	75.17	61.59	*	+
	MAMMA1002938	1.67	1.86	2.42	2.35	3.2	3.56		
	MAMMA1002941	0.49	1.48	2.78	2.53	3.59	2.24		
40	MAMMA1002947	2.24	2.59	4.55	6	6.8	7.94	*	+
	MAMMA1002964	1.73	2.9	5.91	6.91	7.24	7.16	*	+
	MAMMA1002967	1.94	1.59	2.28	2.9	4.19	2.79		
45	MAMMA1002970	2.72	1.77	6	7.59	7.28	8.96	*	+
	MAMMA1002971	1.52	1.6	2.9	2.51	7.27	3.93		
	MAMMA1002972	1	1.32	2.95	1.74	4.56	2.12		
	MAMMA1002973	1.38	2.45	6.73	4.36	6.72	6.78		
50	MAMMA1002979	55.6	60.16	121.72	134.02	101.19	107.19		
	MAMMA1002982	0.53	1.98	2.28	2.04	3.28	1.9		
	MAMMA1002987	1.56	2.11	5.56	3.14	5.55	4.14		
	MAMMA1003003	0.77	2.18	4.78	4.46	6.47	5.08		
55	MAMMA1003004	1.65	1.86	3.7	3.64	3.59	3.16		

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	MAMMA1002712	3.05	3.13	6.98	4.88	7.12	7.39		
	MAMMA1002716	0.56	1.75	3.39	2.38	6.29	2.9		
5	MAMMA1002721	2.11	2.01	-5.57	3.72	6.34	4.59		
	MAMMA1002723	2.43	2.46	4.91	3.85	5.98	4.88		
	MAMMA1002727	3.85	5.55	5.78	5.29	4.45	6.22		
10	MAMMA1002728	21.35	22.03	57.81	49.09	54.73	65.13		
	MAMMA1002742	4.12	4.39	10.35	7.92	8.63	7.61		
	MAMMA1002743	4.12	3.89	6.17	13.81	14.09	13.46	**	+
	MAMMA1002744	2.07	3.15	9.18	9.33	12.98	13.16		
15	MAMMA1002746	0.93	1.28	3.09	2.29	4.31	1.68		
	MAMMA1002748	2.71	2.65	4.52	7.15	5.86	4.72	*	+
	MAMMA1002754	1.12	2.41	5.56	5.05	5.65	6.26		
20	MAMMA1002758	0.71	1.66	2.55	1.57	4.41	1.69		
	MAMMA1002762	11.3	11.14	36.64	38.42	34.23	48.71		
	MAMMA1002764	1.83	3.2	5.95	5.11	6.06	4.26		
	MAMMA1002765	1.19	1.63	4.29	4.63	5.26	2.67		
25	MAMMA1002769	7.4	6.44	13.04	13.78	8.03	12.41		
	MAMMA1002771	1.41	2.41	3.31	3.54	5.39	4.39		
	MAMMA1002775	4.56	4.48	19.79	22.54	29.77	24.29	*	+
30	MAMMA1002780	2.59	1.83	3.03	2.11	4.89	3.78		
	MAMMA1002782	1.43	2.49	3.85	2.51	4.79	4.11		
	MAMMA1002795	1.89	2.03	3.46	6.45	7.68	5.35	**	+
	MAMMA1002796	4.35	3.97	7.51	7.2	8.09	8.17		
35	MAMMA1002805	6.61	11.12	16.52	15.95	24.7	16.5		
	MAMMA1002806	1.47	2.02	3.51	2.28	4.62	2.17		
	MAMMA1002807	1.63	2.4	6.77	6.78	9.66	6.4		
40	MAMMA1002814	3.43	3.52	7.92	9.58	12.39	10.66	*	+
	MAMMA1002817	1.28	1.56	2.87	2.89	5.43	2.91		
	MAMMA1002820	1.66	1.93	2.61	2.52	4.77	2.21		
	MAMMA1002830	67.67	70.46	130.59	165.92	139.33	187.18	*	+
45	MAMMA1002833	4.16	2.88	9.4	8.22	10.68	10.58		
	MAMMA1002835	0.77	1.87	4.03	1.73	3.97	2.79		
	MAMMA1002838	1.85	2.66	5.31	2.91	4.44	3.93		
50	MAMMA1002842	1	3.83	3.84	3.32	4.63	5.15		
	MAMMA1002843	1.72	2.92	2.33	4.09	4.81	3		
	MAMMA1002844	3.05	3.64	6.52	5.26	7.3	4.09		
	MAMMA1002845	1.25	1.57	2.45	3.59	3.55	4.67	*	+
55	MAMMA1002857	92.1	106.97	208.17	209.17	202.29	249.13		
	MAMMA1002858	317.94	188.78	378.89	560.7	620.76	724.33	**	+

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	MAMMA1002585	0.76	1.86	4.38	1.85	3.6	5.26		
	MAMMA1002586	1.98	2.55	3.85	4.12	5.02	3.3		
5	MAMMA1002589	1.08	1.26	-2.44	2.36	5.06	3.19		
	MAMMA1002590	1.01	1.57	5.87	2.58	6.75	4.57		
	MAMMA1002593	2.48	2.48	4.89	4.18	4.07	3.04		
	MAMMA1002597	2.47	2.52	7.25	8.06	9.48	8.78	*	+
10	MAMMA1002598	12.12	13.52	30.83	37.28	48.14	38.91	*	+
	MAMMA1002603	1.2	1.39	3.69	3.25	6.24	4.35		
	MAMMA1002612	3.51	3.39	12.6	7.66	7.78	9.76		
15	MAMMA1002617	4.3	3.41	10.15	6.3	7.29	10.05		
	MAMMA1002618	1.68	2.27	4.02	2.76	3.59	3.91		
	MAMMA1002619	2.96	2.8	5.24	3.22	5.88	3.49		
	MAMMA1002622	2.51	2.12	8.02	7.1	7.18	7.15		
20	MAMMA1002623	2.31	2.21	6.27	5.89	6.17	6.19		
	MAMMA1002625	1.32	1.3	3.23	2.3	6.42	2.6		
	MAMMA1002627	0.98	0.82	2.93	0.6	1.29	0.21		
25	MAMMA1002629	1.8	2.23	6.09	5.03	6.74	7.02		
	MAMMA1002631	1	1.86	3.61	3.07	4.55	2.97		
	MAMMA1002633	6.61	7.44	21.47	19.33	24.55	21.53		
	MAMMA1002636	1.02	2.46	6.97	6.79	8.77	9.25		
30	MAMMA1002637	1.05	1.4	4.66	3.39	4.85	4.28		
	MAMMA1002646	1.69	0.8	3.32	2.33	2.86	1.53		
	MAMMA1002648	10.51	14.07	21.18	42.29	31.45	39.76	**	+
35	MAMMA1002650	1.33	0.56	1.62	1.76	2.08	0.57		
	MAMMA1002652	1.76	2.82	7.31	7.5	7.41	9.79		
	MAMMA1002655	1.7	2.11	3.65	2.54	4.23	3.78		
	MAMMA1002662	0.84	2.24	4.33	3.57	5.68	4.13		
40	MAMMA1002665	3.61	3.57	10.05	13.42	17.97	19.59	*	+
	MAMMA1002671	2.84	3.63	10.17	17.04	16.47	19.3	**	+
	MAMMA1002673	1.32	2.14	4.93	4.07	5.03	2.82		
45	MAMMA1002684	2.95	3.11	3.84	6.61	8.19	7.54	**	+
	MAMMA1002685	0.68	1.49	2.57	2.05	3.74	2.97		
	MAMMA1002692	1.28	1.96	5.45	2.46	4.14	3.62		
	MAMMA1002693	1.84	4.18	8	4.63	7.68	6.61		
50	MAMMA1002698	0.99	1.91	4.05	2.92	4.42	3.3		
	MAMMA1002699	2	2.35	4.43	4.05	5.22	3.64		
	MAMMA1002701	2.41	2.56	8.46	6.72	8.94	8.93		
55	MAMMA1002708	1.51	1.55	5.38	4.08	6.16	6.18		
	MAMMA1002711	1.58	2.08	7.04	4.37	7.35	5.81		

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	MAMMA1002390	1.41	2.04	3.75	5.48	5.4	3.43		
	MAMMA1002392	1.94	3.1	6.1	4.06	5.95	3.32		
5	MAMMA1002396	4.87	3.49	-12.87	10.79	12.9	8.08		
	MAMMA1002399	4.42	5.13	10.69	10.95	8.66	4.57		
	MAMMA1002400	3	2.22	4.69	3.11	4.36	3.53		
10	MAMMA1002409	51.57	55.16	63.3	77.54	80.62	77.88	**	+
	MAMMA1002411	1.08	1.88	4.13	3.43	5.49	1.92		
	MAMMA1002413	2.02	3.01	9.19	5.93	7.17	6.75		
	MAMMA1002417	1.83	2.24	4.87	3.45	4.25	2.63		
15	MAMMA1002427	1.5	2.38	4.54	4.78	5.56	3.41		
	MAMMA1002428	2.47	2.26	5.38	4.46	5.11	4.28		
	MAMMA1002433	1.74	2.18	6.84	6.72	6.96	6.22		
20	MAMMA1002434	2.94	2.4	7.38	5.34	4.65	5.03		
	MAMMA1002446	1.39	2.34	5.62	3.98	5.84	5.96		
	MAMMA1002447	2.51	1.38	6.4	5.11	6.26	5.45		
	MAMMA1002454	7.77	9.16	18.07	21.71	17.12	18.35		
25	MAMMA1002461	2.06	4.11	7.7	4.92	5.41	6.47		
	MAMMA1002463	3.28	3.32	8.09	6.98	7.82	5.39		
	MAMMA1002464	16.58	16.77	20.05	19.41	20.41	18.09		
30	MAMMA1002466	9.48	9.89	14.22	14.58	15.75	13.93		
	MAMMA1002470	1.39	1.51	5.13	3.54	5.01	3.73		
	MAMMA1002475	0.72	1.85	5.03	3.86	5.17	4.65		
	MAMMA1002480	0.66	1.21	2.31	1.68	2.84	2.03		
35	MAMMA1002485	29.98	27.24	46.09	64.83	74.9	80.68	**	+
	MAMMA1002494	2	2	4.11	4.48	5.12	5.13	*	+
	MAMMA1002498	0.97	2.57	3.16	2.07	3.18	1.55		
40	MAMMA1002524	3.04	2.96	6.43	5.18	7.34	6.1		
	MAMMA1002530	2.5	3.24	4.88	3.17	4.41	2.55		
	MAMMA1002538	2.34	2.38	5.62	5.46	5.13	4.91		
	MAMMA1002545	2.37	2.64	6.26	4.56	6.49	4.56		
45	MAMMA1002554	1.96	1.42	5.43	5.3	6.01	7.81		
	MAMMA1002556	1.3	1.9	3.6	3.73	5.75	3.89		
	MAMMA1002561	2.3	2.99	7.19	8.13	10.46	7.98		
50	MAMMA1002565	1.22	2.15	3.52	2.57	4.51	2.55		
	MAMMA1002566	0.98	1.87	6.21	1.65	4.7	3.9		
	MAMMA1002571	0.53	1.8	3.06	1.43	3.1	4.3		
	MAMMA1002573	2.14	1.86	7.06	4.54	5.66	5.97		
55	MAMMA1002576	118.77	131.84	363.97	348.62	471.73	358.66		
	MAMMA1002584	3.52	2.27	11.91	12.86	17.82	13.46		

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	MAMMA1002250	1.06	2.09	5.2	3.95	6.82	6.01		
	MAMMA1002253	2.77	2.39	3.45	4.84	6.18	3.37		
5	MAMMA1002267	17.17	19.95	-51.7	130.02	108.53	115.75	**	+
	MAMMA1002268	1.72	2.28	5.82	6.92	11.3	6.52		
	MAMMA1002269	0.89	0.73	2.25	2.32	2.58	1.67		
	MAMMA1002282	0.86	1.09	4.95	5.87	5.31	6.81		
10	MAMMA1002292	2.71	2.25	7.77	10.57	10.52	11.53	*	+
	MAMMA1002293	3.71	3.31	12.81	8.54	10.47	12.05		
	MAMMA1002294	0.9	1.71	4.61	3.68	6.03	4.2		
15	MAMMA1002297	1.53	3.25	7.45	5.77	7.8	6.91		
	MAMMA1002298	1.48	1.4	3.98	3.85	3.11	2.46		
	MAMMA1002299	1.5	1.69	3.16	3.91	2.97	2.2		
	MAMMA1002308	1.39	1.35	6.55	4.5	3.11	2.54		
20	MAMMA1002310	3.56	3.84	12.73	9.92	12.66	11.48		
	MAMMA1002311	2.52	2.13	6.82	9.61	9.66	6.9	*	+
	MAMMA1002312	1.63	2.22	5.19	3.51	8.45	2.55		
25	MAMMA1002317	2.08	2.55	4.89	4.08	3.85	4.09		
	MAMMA1002319	0.8	2.78	3.51	2.68	3.97	2.85		
	MAMMA1002322	2.48	3.23	7.84	12.21	10.02	8.55	*	+
	MAMMA1002329	1.64	1.67	2.93	2.9	3.3	2.76		
30	MAMMA1002332	2.17	2.38	4.58	5.98	4.14	3.05		
	MAMMA1002333	1.7	1.74	4.19	5.35	5.07	3.54		
	MAMMA1002335	1.75	2.72	8.53	6.93	11.32	4.23		
35	MAMMA1002339	2.09	2.42	7.34	5.21	7.5	5.14		
	MAMMA1002347	1.7	2.3	6.39	5.5	5.32	4.64		
	MAMMA1002351	2.08	2.68	5.74	3.03	4.48	4.84		
	MAMMA1002352	1.27	2.28	3.66	3.53	4.63	2.8		
40	MAMMA1002353	4.46	2.5	5.84	5.95	4.19	4		
	MAMMA1002355	3.97	3.38	8.37	7.98	7.31	8.57		
	MAMMA1002356	2.18	1.49	4.36	5.43	4.13	3.75		
45	MAMMA1002359	3.95	3.35	16.09	23.81	24.53	19	*	+
	MAMMA1002360	0.93	1.73	3.77	2.48	3.2	1.67		
	MAMMA1002361	2.01	2.64	4.53	4.17	4.95	4.03		
	MAMMA1002362	2.33	2.33	3.36	5.31	5.51	3.99	*	+
50	MAMMA1002367	2.97	3.64	14.63	18.34	21.06	21.56	*	+
	MAMMA1002371	2.28	3.75	8.3	6.15	6.74	5.88		
	MAMMA1002380	1.81	2.26	4.9	4.71	5.76	3.55		
	MAMMA1002384	2.14	1.53	4.73	4.48	5.36	4.05		
55	MAMMA1002385	1.19	2.05	5.63	3.34	4.8	2.47		

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	MAMMA1002091	3.79	3.22	4.32	7.18	6.76	6.41	**	+
	MAMMA1002093	0.72	1.4	4.31	2.74	4.39	2.33		
5	MAMMA1002095	2.4	3.22	7.5	4.73	7.52	4.54		
	MAMMA1002108	1.84	1.02	2.63	1.87	3.24	1.31		
	MAMMA1002112	2.94	3.4	7.03	12.79	16.02	11.28	**	+
10	MAMMA1002118	1.02	1.61	2.24	1.41	3.18	2.01		
	MAMMA1002119	0.76	2.15	3.61	1.51	3.12	2.54		
	MAMMA1002125	1.79	2.61	6.95	4.52	4.19	4.11		
	MAMMA1002126	3.72	4.25	9.79	10.08	9.02	11.03		
15	MAMMA1002128	0.9	2.36	3.07	2.7	3.49	2.88		
	MAMMA1002132	3.78	3.24	11.42	6.18	9.05	6.81		
	MAMMA1002140	1.46	1.87	3.68	2.18	3.24	2.33		
20	MAMMA1002142	3.13	3.43	7.06	5.18	7.62	5.46		
	MAMMA1002143	5.42	2.27	7.96	7.98	9.87	13.23		
	MAMMA1002145	1.47	1.34	3.3	2.9	4.02	2.64		
	MAMMA1002147	0.81	1.59	2.9	2.71	4.4	3		
25	MAMMA1002153	0.99	1.92	5.55	3.52	6.41	4.75		
	MAMMA1002155	2.11	1.93	6.76	4.4	6.46	4.46		
	MAMMA1002156	0.81	0.8	1.94	0.67	2.63	0.78		
30	MAMMA1002158	1.38	1.83	5.12	4.09	7.73	5.2		
	MAMMA1002164	2.01	2.09	5.86	3.17	3.18	4.04		
	MAMMA1002165	4.04	4.29	7.25	8.65	8.1	6.81		
	MAMMA1002170	1.01	1.48	154.53	2.65	3.24	4.11		
35	MAMMA1002174	1.66	2.9	5.88	4.55	7.78	8.58		
	MAMMA1002175	3.27	3.3	7.02	6.95	6.64	7.22		
	MAMMA1002180	8.59	6.53	35.97	55.49	48.49	51.08	*	+
40	MAMMA1002198	3.11	2.3	9.33	7.6	11.22	7.13		
	MAMMA1002205	2.93	1.66	6.15	6.3	8.04	7.54		
	MAMMA1002206	4.6	3.59	8.14	12.4	13.97	11.74	**	+
	MAMMA1002209	1.7	1.93	4.03	4.43	4.23	4.57		
45	MAMMA1002215	4.17	2.72	15.2	11.05	12.43	17.14		
	MAMMA1002219	1.57	1.96	4.99	4.84	6.34	5.96		
	MAMMA1002224	3.18	2.9	8.18	5.49	7.25	5.86		
50	MAMMA1002229	3.74	2.21	8.83	8.48	9.26	6.82		
	MAMMA1002230	2.02	2.21	6.63	5.31	8.91	6.58		
	MAMMA1002233	3.01	1.6	6.08	4.21	7.91	6.14		
	MAMMA1002234	3.05	3.06	6.7	8.6	10.45	10.76	*	+
55	MAMMA1002236	4.13	3.68	14.08	26.56	20.38	24.71	*	+
	MAMMA1002243	0.97	2.48	3.48	3.28	3.43	2.96		

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	MAMMA1001893	3.74	3.42	6.25	6.59	5.49	5.58		
	MAMMA1001901	1.13	1.5	5.4	4.53	5.72	2.67		
5	MAMMA1001907	2.57	1.62	6.43	4.15	7.36	6.34		
	MAMMA1001908	3.2	3.36	8.35	11.83	12.96	12.46	*	+
	MAMMA1001919	0.23	0.97	3.3	2.24	3.9	2.07		
10	MAMMA1001931	0.76	1.65	4.04	3.36	5.89	3.25		
	MAMMA1001937	2.27	3.15	5.5	6.44	5.06	3.78		
	MAMMA1001951	1.74	2.57	6.47	6.48	6.15	4.83		
	MAMMA1001956	3.02	3.48	9.72	8.52	7.66	6.76		
15	MAMMA1001957	3.39	3.51	9.15	7.88	9.47	7.66		
	MAMMA1001960	3.1	3.34	7.24	12.06	9.14	6.1		
	MAMMA1001963	0.57	0.78	2.14	1.3	2.36	1.06		
20	MAMMA1001969	1.7	3.43	10.86	8.54	11.14	8.74		
	MAMMA1001970	2.86	3.04	8.48	13.11	6.59	6.64		
	MAMMA1001978	0.57	1.85	1.76	2.42	3.87	1.53		
	MAMMA1001992	2.07	2.04	5.65	6.79	6.75	5.09		
25	MAMMA1001994	7.97	3.65	11	18.83	13.23	17.17	*	+
	MAMMA1002008	3.28	3.77	6.42	3.43	4.06	1.24		
	MAMMA1002009	1.46	2.94	5.17	5.73	7.57	4.06		
30	MAMMA1002011	1.77	1.71	4.26	6.5	6.45	3.37		
	MAMMA1002022	1.51	2.1	5.92	6.64	7.42	5.2		
	MAMMA1002024	9.79	9.67	19.03	17.61	16.96	22.43		
	MAMMA1002032	2.78	2.41	7.25	5.29	6.16	8.07		
35	MAMMA1002033	3.23	3.95	7.73	11.24	7.23	6.62		
	MAMMA1002041	2.87	2.25	3.18	4.74	5.39	1.71		
	MAMMA1002042	2.54	2.34	5.66	5.65	5.78	3.76		
40	MAMMA1002045	2.33	3.51	7.28	8.39	5.05	4.44		
	MAMMA1002047	2.58	2.98	8.83	8.7	8.9	6.89		
	MAMMA1002056	2.01	5.78	11.14	11.35	10.64	9.14		
	MAMMA1002058	1.67	2.61	8.19	4.84	4.66	4.27		
45	MAMMA1002060	1.08	2.08	1.41	2.5	4.09	1.2		
	MAMMA1002065	1.81	2.75	6.04	7.19	5.19	3.26		
	MAMMA1002068	2.43	1.84	5.29	4.98	5.6	4.47		
	MAMMA1002070	4.5	2.92	4.15	2.58	5.23	2.81		
50	MAMMA1002078	1.32	1.43	2.94	1.12	4.4	1.07		
	MAMMA1002080	7.98	9.71	13.38	14.92	20.84	14.26		
	MAMMA1002082	2.54	4.96	13.04	9.67	8.15	7.78		
55	MAMMA1002084	1.78	3.47	3.38	4.68	4.48	3.6		
	MAMMA1002087	1.12	2.15	5.37	3.6	4.67	2.36		

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	MAMMA1001757	1.21	1.1	2.32	2.21	3.25	2.43		
	MAMMA1001760	3.87	4.52	20.01	22.91	24.2	27.59	*	+
5	MAMMA1001764	2.62	2.36	5.97	7.13	10.17	6.51		
	MAMMA1001767	1.22	1.55	2.13	1.61	2.96	1.55		
	MAMMA1001768	0.57	1.18	4.25	4.74	4.72	4.37		
10	MAMMA1001769	2.48	2.83	9.22	9.3	9.81	8.94		
	MAMMA1001771	2.66	1.58	3.74	2.86	5.85	6.77		
	MAMMA1001773	2.7	3.53	3.87	4	6.29	7.61		
	MAMMA1001778	0.88	1.92	3.14	3.13	4.21	3.61		
15	MAMMA1001783	2.01	2.1	11.25	11.63	18.46	13.04		
	MAMMA1001785	3	3.52	8.85	10.56	13.38	11	*	+
	MAMMA1001788	0.49	0.86	1.21	0.72	1.72	1.11		
20	MAMMA1001790	1.68	1.67	5.1	2.37	3.73	3.93		
	MAMMA1001800	0.83	0.99	1.47	1.5	2.24	3.25		
	MAMMA1001804	1.02	1.41	3.18	2.37	4.16	2.4		
	MAMMA1001806	2.13	2.78	6.4	3.15	5.5	4.72		
25	MAMMA1001812	1.46	1.33	5.52	4.21	5.86	5.05		
	MAMMA1001815	0.33	1.76	3.07	1.22	3.67	1.24		
	MAMMA1001817	3.19	3.38	9.5	6.78	10.89	13.3		
30	MAMMA1001818	1.68	2.08	3.41	3.94	8.52	3.41		
	MAMMA1001819	2.57	4.12	5.82	8.7	10.29	7.87	*	+
	MAMMA1001820	2.68	4.51	8.27	7.51	10.98	6.07		
	MAMMA1001824	1.66	2.83	8.36	7.55	9.8	7.11		
35	MAMMA1001832	6.72	7.99	11.85	20.17	21.28	17.21	**	+
	MAMMA1001836	1.74	1.66	5.08	4.79	8.19	4.88		
	MAMMA1001837	2.61	2	5.84	7.1	9.19	5.37		
40	MAMMA1001848	1.02	1.61	3.3	2.81	5.33	3.18		
	MAMMA1001850	3.79	4.51	9.31	9.98	9.93	14.19		
	MAMMA1001851	1.49	2.33	4.98	4.97	4.12	4.02		
	MAMMA1001852	2.98	4	9.68	6.4	7.56	6.8		
45	MAMMA1001854	2.56	3.11	9.16	10.59	10.64	9.98		
	MAMMA1001858	3.11	2.22	5.28	9.93	7.91	8.87	**	+
	MAMMA1001864	1.69	1.91	4.09	8.91	6.18	4.37		
50	MAMMA1001868	0.71	0.92	2.64	1.68	2.58	0.91		
	MAMMA1001874	1.2	0.87	2.52	1.06	3.48	1.17		
	MAMMA1001878	3.1	3.46	10.86	7.7	13.37	6.77		
	MAMMA1001880	2.67	2.99	7.24	5.58	7.17	8.12		
55	MAMMA1001885	1.14	1.93	6.19	4.7	5.54	4.58		
	MAMMA1001890	3.54	3.95	12.93	13.59	13.29	12.2		

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	MAMMA1001599	4.64	7.15	16.79	15.8	15.18	15.06		
	MAMMA1001600	1.45	2.22	4.73	2.98	4.68	2.11		
5	MAMMA1001604	1.03	1.76	3.62	2.35	4.01	1.64		
	MAMMA1001606	1.64	2.04	5.15	3.58	5.45	4.27		
	MAMMA1001609	1.31	2.37	4.36	3.05	5.43	1.59		
	MAMMA1001614	2.91	3.57	6.15	5.94	6.11	4.14		
10	MAMMA1001615	3.98	2.61	10.12	8.87	8.29	8.41		
	MAMMA1001619	7.73	7.8	14.29	16.33	12.93	14.61		
	MAMMA1001620	2.53	2.41	7.98	5.77	7.13	4.54		
15	MAMMA1001623	4.11	4.58	9.3	7.34	9.28	6.75		
	MAMMA1001626	0.83	1.98	2.52	3.24	3.93	1.93		
	MAMMA1001627	1.11	1.98	3.57	2.63	3.68	1.63		
	MAMMA1001630	2.02	3.08	7.83	7.49	7.53	4.29		
20	MAMMA1001633	2.64	3.12	8.8	12.09	9.59	6.16		
	MAMMA1001634	2.83	2.7	6.11	8.69	8.27	6.9	*	+
	MAMMA1001635	5.65	2.39	9.52	7.92	8.3	8.37		
25	MAMMA1001649	1.61	1.63	4.71	2.95	4.62	2.53		
	MAMMA1001654	8.14	9.45	39	43.4	55	46.79		
	MAMMA1001660	19.61	17.92	37.43	40.94	27.03	34		
	MAMMA1001663	1.9	4.73	9.42	9.59	9.56	6.58		
30	MAMMA1001670	1.12	2.66	3.97	3.65	4.09	2.62		
	MAMMA1001671	1.08	1.42	3.56	1.37	4.64	1.77		
	MAMMA1001679	6.85	6.37	13.89	11.48	17.04	13.91		
35	MAMMA1001683	2.15	3.29	9.6	6.58	6.53	6.96		
	MAMMA1001686	1.25	1.34	3.77	1.39	2.97	3.06		
	MAMMA1001688	113.39	113.61	245.56	392.2	458.41	413	**	+
	MAMMA1001689	1.01	3.76	4.1	5.04	3.79	4.44		
40	MAMMA1001692	1.97	2.59	5.37	3.66	5.3	3.88		
	MAMMA1001711	1.99	3.64	8.65	4.35	5.51	6.1		
	MAMMA1001715	1.31	1.64	3.95	4.64	4.87	4.13		
45	MAMMA1001730	2.01	2.15	2.5	2.8	4.42	2.83		
	MAMMA1001735	44.73	48.32	102.35	94.99	156.23	119.88		
	MAMMA1001740	0.64	1.6	4.59	2.06	3.91	1.95		
	MAMMA1001743	9.84	11.15	33.16	41.97	51.62	49.6	*	+
50	MAMMA1001744	0.63	0.72	0.86	1.1	1.72	1.71	*	+
	MAMMA1001745	1.41	2.15	6.15	3.27	4.46	3.93		
	MAMMA1001751	1.38	2.41	3.24	2.85	4.51	4.32		
	MAMMA1001752	4.7	4.78	9.75	6.12	9.61	8.4		
55	MAMMA1001754	7.25	7.89	7.34	11.04	9.63	9.39	*	+

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	MAMMA1001401	26.87	16.48	32.72	43.47	57.55	45.66	*	+
	MAMMA1001408	1.06	1.06	2.57	0.65	4.22	1.19		
5	MAMMA1001411	1.65	1.26	3.84	4.38	3.33	3.51		
	MAMMA1001414	3.12	3.85	5.74	12.58	10.67	15.28	**	+
	MAMMA1001415	2.45	3.16	11.93	14.57	20.15	13.69	*	+
10	MAMMA1001418	0.66	2.2	5.36	3.57	6.04	4.46		
	MAMMA1001419	0.8	2.43	4.93	6.03	7.01	3.92		
	MAMMA1001420	0.96	3.09	4.5	3.23	4.11	3.41		
	MAMMA1001426	20.24	32.21	42.42	44.31	39.63	38.75		
15	MAMMA1001428	1.94	2.83	6.35	3.8	6.93	4.33		
	MAMMA1001432	1.19	2.33	8.19	5.62	6.19	6.68		
	MAMMA1001435	1.43	0.78	3.32	3.48	3.67	2.64		
20	MAMMA1001442	1.96	3.94	7.41	8.18	8.6	6.63		
	MAMMA1001446	2.17	2.57	6.71	6.9	7.34	7.97		
	MAMMA1001450	1.22	2.05	3.58	2.81	4.18	2.39		
	MAMMA1001452	1.99	1.78	5.92	8.38	6.19	4.83		
25	MAMMA1001465	3.93	3.25	13.61	16.65	14.6	13.82		
	MAMMA1001476	1.63	1.09	4.25	5.87	5.95	4.64	*	+
	MAMMA1001478	2.28	2.12	5.98	3.55	6.27	4.19		
30	MAMMA1001479	3.11	4.71	8.32	5.58	6.74	6.21		
	MAMMA1001487	1.1	1.14	3.84	4.73	3.26	2.08		
	MAMMA1001498	1.93	3.41	7.78	6.17	7.45	5.64		
	MAMMA1001501	0.88	1.97	4.49	2.8	4.77	2.36		
35	MAMMA1001502	1.82	1.91	6.48	3.29	6.29	6.26		
	MAMMA1001510	0.48	0.78	2.92	0.54	3.04	1.19		
	MAMMA1001522	1.03	1.29	3.94	5.05	4.9	3.39		
40	MAMMA1001529	0.72	2.06	3.22	3.74	4.07	2.57		
	MAMMA1001532	1.74	1.86	4.27	3.79	5.71	3.12		
	MAMMA1001533	0.61	1.31	2.9	1.52	3.06	1.64		
	MAMMA1001534	0.44	2.59	2.4	1.48	3.64	1.14		
45	MAMMA1001535	1.38	1.91	3.99	2.12	3.98	2.38		
	MAMMA1001547	2.8	2.89	7.77	9.23	8.22	6.22		
	MAMMA1001551	1.1	1.48	4.46	2.23	2.88	2.99		
50	MAMMA1001569	1.27	1.68	3.41	2.03	3.41	1.94		
	MAMMA1001575	1.48	2.41	3.42	4.01	4.43	2.81		
	MAMMA1001576	4.79	8.23	9.65	14.75	9.39	17.03		
	MAMMA1001584	0.89	2.48	3.33	3.11	4	3.09		
55	MAMMA1001586	1.43	2.41	3.34	3.78	3.31	1.84		
	MAMMA1001590	2.96	2.53	5.55	5.44	6.47	6.04		

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	MAMMA1001223	2.48	3.32	6.53	4.95	6.51	4.1		
	MAMMA1001232	2.82	4.27	8.08	12.22	8.82	9.57		
5	MAMMA1001234	1.25	3.27	-3.17	5.05	3.91	3.26		
	MAMMA1001237	1.22	1.56	4.21	1.94	3.66	2.09		
	MAMMA1001243	2.18	2.28	4.06	4.05	4.89	1.99		
	MAMMA1001244	1.22	1.16	2.86	2.96	4.79	2.22		
10	MAMMA1001249	2.3	1.89	5.93	5.19	5.8	3.75		
	MAMMA1001256	3	3.09	8.29	5.89	7.83	8.01		
	MAMMA1001259	4.38	3.25	7.15	7.94	9.24	6.63		
15	MAMMA1001260	1.76	2.71	5.42	6.51	5.33	7.33		
	MAMMA1001262	2.1	4.11	5.28	7.86	8.04	6.25	*	+
	MAMMA1001268	2	2.16	4.59	2.56	4.23	2.48		
	MAMMA1001271	4.84	5.78	17.37	18.29	14.24	15.67		
20	MAMMA1001274	2.88	3.06	6.17	6.22	8.55	7.93		
	MAMMA1001280	2.09	1.48	4.36	1.84	3.78	1.73		
	MAMMA1001283	1.63	1.71	6.34	6.88	5.63	4.83		
25	MAMMA1001284	2.27	2	8.67	5.08	9.09	9.51		
	MAMMA1001286	13.83	9.72	17.39	12.15	11.83	14.63		
	MAMMA1001289	17.63	13.49	23.32	21.02	26.39	36.8		
	MAMMA1001292	3	3.01	5.94	7.26	6.31	6.85	*	+
30	MAMMA1001296	3.55	3.76	12.61	14.11	12.37	12.8		
	MAMMA1001298	1.26	1.7	6.26	4.25	6.78	4.07		
	MAMMA1001305	0.86	1.59	4.43	2.49	4.07	2.63		
35	MAMMA1001309	0.61	0.9	2.7	1.84	3	1.49		
	MAMMA1001310	1.72	2.17	3.64	4.81	7.38	4.42		
	MAMMA1001322	0.99	1.54	1.83	2.83	1.77	2.13		
	MAMMA1001324	1.3	1.12	3.16	2.03	2.83	1.94		
40	MAMMA1001330	3.35	2.65	9.53	7.93	9.75	5.36		
	MAMMA1001333	3.1	3.74	10.23	9.88	11.4	9.07		
	MAMMA1001334	5.53	4.17	4.83	10.97	8.23	10.16	**	+
45	MAMMA1001337	2.49	3.54	6.6	6.99	9.16	8.05	*	+
	MAMMA1001341	1.21	1.14	3.48	1.54	5.66	1.41		
	MAMMA1001343	2.37	1.89	8.07	8.17	9.75	10.95		
	MAMMA1001344	9.59	9.07	11.75	13.63	11.67	15.98		
50	MAMMA1001346	1.34	1.25	3.9	2.05	3.9	2.94		
	MAMMA1001383	3.07	3.61	8.52	8.3	9.02	9.38		
	MAMMA1001388	1.62	1.93	5.34	3.38	6.11	4.58		
55	MAMMA1001396	4.2	2.12	8.12	11.39	10.42	8.68		
	MAMMA1001397	2.59	2.27	5.79	8.33	8.96	7.78	*	+

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	MAMMA1001054	2.04	2.58	6.99	9.29	11.07	8.36	*	+
	MAMMA1001059	2.66	4.71	9.73	8.85	9.72	8.46		
5	MAMMA1001066	3.64	2.97	12.26	14.08	12.09	8.96		
	MAMMA1001067	1.26	2	4.77	3.53	5.33	3.29		
	MAMMA1001072	1.44	2.06	7.76	6.38	7.13	7.2		
	MAMMA1001073	1.17	0.79	1.47	1.49	2.74	1.34		
10	MAMMA1001074	0.78	1.47	3.97	6.24	5.4	4.9	*	+
	MAMMA1001075	4.87	4.41	10.48	11.24	9.82	8.21		
	MAMMA1001078	1.7	1.83	7.96	9.69	9.99	9.37	*	+
15	MAMMA1001080	3.77	3.93	6.97	8.71	10.33	5.57		
	MAMMA1001082	1.51	2.03	4.03	1.73	5.23	1.6		
	MAMMA1001091	1.17	1.36	2.02	1.81	3.03	1.47		
	MAMMA1001092	1.93	2.09	4.81	3.17	3.57	1.89		
20	MAMMA1001094	1.73	4.28	4.65	4	5.62	4.7		
	MAMMA1001105	2.45	2.62	7.7	6.99	7.57	6.06		
	MAMMA1001110	0.4	1.01	2.74	1.42	2.53	0.47		
25	MAMMA1001126	1.96	3.09	10.92	8.39	9.27	6.08		
	MAMMA1001133	2.5	3.44	10.94	9.48	10.83	10.68		
	MAMMA1001139	87.88	86.18	214.31	193.19	47.2	160.92		
	MAMMA1001141	1.33	2.89	3.65	3.69	5.25	5.46		
30	MAMMA1001143	2.02	1.79	4.23	3.95	6.69	4.34		
	MAMMA1001145	3.1	2.22	3.39	6.37	7.13	2.84		
	MAMMA1001150	1.34	2.48	3.95	3.06	3.31	2.11		
35	MAMMA1001154	2.16	2.8	5.57	5.44	7.13	5.22		
	MAMMA1001159	4.19	4.01	11.06	11.31	5.89	9.45		
	MAMMA1001161	4.3	5.27	19.53	18.34	10.8	14.8		
	MAMMA1001162	1.98	1.77	3.16	5.25	5.13	2.25		
40	MAMMA1001181	2.44	2.28	4.87	5.06	4.74	3.62		
	MAMMA1001186	2	2.66	4.66	5.38	5.48	3.9		
	MAMMA1001189	2.23	3.68	7.17	11	11.17	9.9	*	+
45	MAMMA1001191	2.54	2.07	5.49	4.37	3.89	2.97		
	MAMMA1001198	368.47	416.05	784.82	647.17	738.61	605.52		
	MAMMA1001202	11.78	11.85	30.06	34.39	28.74	25.16		
	MAMMA1001203	2.57	3.01	7.15	8.72	6.26	5.56		
50	MAMMA1001206	1.91	3.28	4.5	3.69	6.66	2.65		
	MAMMA1001208	2.66	2.93	3.31	3.82	4.95	3.19		
	MAMMA1001215	2.9	3.08	6.55	3.49	8.09	4.74		
55	MAMMA1001220	2.63	3.03	7.25	7.16	7.17	6.03		
	MAMMA1001222	1.25	1.18	4.18	2.18	5.85	0.53		

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	MAMMA1000921	1.03	1.02	2.41	3.47	3.84	3.17	*	+
	MAMMA1000931	2.68	3.44	3.95	6.78	7.66	7.32	**	+
5	MAMMA1000940	1.67	1.84	6.05	5.78	6.84	6.3		
	MAMMA1000941	3.74	2.55	8.61	9.01	10.11	9.46		
	MAMMA1000942	2.75	1.85	7.46	8.27	8.47	7.2		
	MAMMA1000943	2.16	2.84	10.49	9.04	11.08	8.74		
10	MAMMA1000952	2.6	1.93	8.65	9.03	7.47	6.75		
	MAMMA1000956	0.93	1.24	3.11	3.64	3.1	3.47		
	MAMMA1000957	2.5	1.41	2.62	4.81	5.73	6.85	**	+
15	MAMMA1000962	3.25	3.57	10.48	13.62	11.18	16.85	*	+
	MAMMA1000966	1.85	2.19	6.04	6.34	6.23	7.04		
	MAMMA1000968	1.6	1.46	5.49	4.79	5.62	4.97		
	MAMMA1000972	2.4	1.41	3.83	3.34	3.91	4.32		
20	MAMMA1000973	5.14	3.37	12.58	7.02	8.56	10.31		
	MAMMA1000975	1.44	1.99	3.34	2.37	4.9	5.05		
	MAMMA1000976	2.46	2.71	8.57	9.22	11.17	8.92		
25	MAMMA1000979	1.46	2.62	3.06	4.34	4.41	7.71		
	MAMMA1000986	5.75	5.32	10.24	8.83	11.32	13.95		
	MAMMA1000987	1.44	1.36	3.99	2.43	3.74	5.66		
	MAMMA1000988	3.76	4.86	8.88	10.18	11.34	10.5	*	+
30	MAMMA1000994	9.82	7.58	15.88	12.02	11.56	9.25		
	MAMMA1000998	1.51	1.07	3.13	4.21	5.42	4.04	*	+
	MAMMA1001003	1.98	1.83	5.97	4.2	6.86	4.39		
35	MAMMA1001007	0.38	1.03	1.77	0.14	1.38	0.32		
	MAMMA1001008	11.76	11.09	40.52	56.73	50.93	45.37	*	+
	MAMMA1001013	3.62	4.16	12.14	8.42	11.54	10.08		
	MAMMA1001014	1.4	1.79	5.49	4.36	5.16	4.26		
40	MAMMA1001021	0.49	2.08	7.85	6.46	5.47	4.26		
	MAMMA1001024	0.85	1.59	3.14	2.55	3.92	1.96		
	MAMMA1001025	1.03	1.47	2.94	1.27	2.95	1.41		
45	MAMMA1001028	1.3	1.09	2.23	3.07	4.73	3.28	*	+
	MAMMA1001030	1.63	0.48	2.22	3.06	3.24	2.08		
	MAMMA1001035	2.48	2.92	10.31	9.29	11.92	11.54		
	MAMMA1001036	4.69	4.01	10.7	11.23	8.22	11.33		
50	MAMMA1001037	1.91	2.88	6.49	5.28	7.53	4.69		
	MAMMA1001038	1.18	1.59	4.28	4.12	3.85	4.45		
	MAMMA1001041	1.64	1.87	3.18	3.75	4.04	2.64		
55	MAMMA1001043	1.09	1.24	3.67	3.44	3.35	3.18		
	MAMMA1001050	1.52	1.55	5.94	8.4	8.28	6.51	*	+

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	MAMMA1000788	3.05	4.31	4.38	4.16	5.12	2.21		
	MAMMA1000798	1.01	2.86	2.77	2.03	3.56	2.1		
5	MAMMA1000802	4.36	3.71	8.23	17.49	15.39	19.2	**	+
	MAMMA1000810	3.91	4.98	14.15	17.58	19.04	16.15	*	+
	MAMMA1000813	1.63	2.11	3.73	2.82	3.15	2.06		
	MAMMA1000814	2.56	2.97	7.82	7.28	7.26	6.61		
10	MAMMA1000824	12.58	11.27	34.16	62.44	72.28	50.62	*	+
	MAMMA1000827	1.83	2.04	5.05	3.3	4.77	4.31		
	MAMMA1000831	1.45	2.81	2.84	2.55	5.43	1.85		
15	MAMMA1000838	6.85	9.27	7.86	13.01	8.43	14.52		
	MAMMA1000839	4.89	4.41	9	9.02	8.66	14.04		
	MAMMA1000841	1.33	2.02	3.11	2.23	4.44	2.29		
20	MAMMA1000842	2.48	1.88	3.59	3.03	3.74	1.82		
	MAMMA1000843	1.26	2.19	3.88	2.26	4.43	2.35		
	MAMMA1000845	0.83	1.01	2.4	2.35	3.88	1.24		
	MAMMA1000851	1.3	3.42	5.35	7.83	5.9	6.65		
25	MAMMA1000854	2.77	3.7	6.33	5.08	5.68	5.13		
	MAMMA1000855	0.37	2.97	2.62	2.51	3.74	1.78		
	MAMMA1000856	0.87	1.39	3.11	2.05	6.37	3.19		
	MAMMA1000859	9.88	8.56	20.5	19.52	18.47	24.31		
30	MAMMA1000862	1.13	1.55	3.53	1.17	3.18	0.79		
	MAMMA1000863	2.62	2.08	4.72	2.59	5.24	4.91		
	MAMMA1000865	0.35	0.82	2.48	0.4	1.84	0.35		
35	MAMMA1000867	1.08	2.83	2.87	1.95	3.68	4.76		
	MAMMA1000875	0.89	2.72	2.34	3.31	3.57	2.59		
	MAMMA1000876	1.23	1.64	4.59	2.4	3.37	3.22		
	MAMMA1000877	3.15	2.89	9.22	8.39	10.32	10.07		
40	MAMMA1000878	3.05	3.61	8.94	6.33	8.52	9.62		
	MAMMA1000880	1.46	1.15	4.77	2.45	3.96	3.01		
	MAMMA1000881	1.81	2.09	4.52	3.77	6.07	4.73		
45	MAMMA1000883	0.57	0.79	2.1	1.37	2.14	2.11		
	MAMMA1000897	0.76	2.39	0.36	1.19	2.99	6.19		
	MAMMA1000898	1.06	1.99	1.75	1.41	2.49	3.7		
	MAMMA1000905	1.8	2.75	4.68	8.32	6.86	10.64	*	+
50	MAMMA1000906	1.17	2.49	2.63	2.45	4.27	4.08		
	MAMMA1000908	1.59	1.63	4.3	1.77	3.05	1.43		
	MAMMA1000911	4.97	6.25	8.37	21.77	20.01	20.98	**	+
55	MAMMA1000914	1.14	0.85	2.41	1.1	2.23	1.61		
	MAMMA1000920	1.99	2.17	4.41	10.82	10.67	9.11	**	+

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	MAMMA1000652	1.98	1.61	3.56	4.21	5.34	5.24	*	+
	MAMMA1000657	2.28	1.78	4.51	2.18	4.48	3.26		
5	MAMMA1000664	1.78	1.49	5.35	2.9	6.43	6.85		
	MAMMA1000667	1.24	1.68	2.17	1.96	5.64	2.41		
	MAMMA1000668	0.71	1.11	3.23	2.44	3.76	1.74		
	MAMMA1000669	0.76	0.97	2.01	0.9	3.94	2.06		
10	MAMMA1000670	3.27	2.78	4.47	10.4	6.73	9.28	*	+
	MAMMA1000672	1.71	3.23	6.88	5.43	5.63	6.03		
	MAMMA1000681	0.98	1.19	2.53	1.98	3.73	2.45		
15	MAMMA1000684	6.87	11.61	18.54	30.76	32.53	30.62	**	+
	MAMMA1000696	1.64	3.39	4.99	7.89	14.39	8.69	*	+
	MAMMA1000702	3.12	3.07	5.9	7.47	10.05	7.55	*	+
	MAMMA1000706	0.63	1.07	1.79	1.08	1.52	0.66		
20	MAMMA1000707	0.74	1.26	1.76	0.83	1.87	0.63		
	MAMMA1000713	1.53	2.14	5.33	5.43	5.8	6.96		
	MAMMA1000714	1.19	1.84	4.31	2.64	4.96	4.94		
25	MAMMA1000718	1.32	2.79	4.84	5.53	7.12	4.37		
	MAMMA1000720	1.33	2.19	5.14	4.95	8.51	5.44		
	MAMMA1000723	1.22	1.65	4.17	3.26	4.81	3.68		
	MAMMA1000731	1.24	1.17	3.11	3.04	4.99	3.26		
30	MAMMA1000732	1.37	1.59	3.02	4.86	6.5	6.05	**	+
	MAMMA1000733	0.58	0.82	1.54	2.31	2.41	1.22		
	MAMMA1000734	12.22	11.56	22.62	21.95	22.19	13.18		
35	MAMMA1000736	4.26	4.34	11.96	4.92	5.77	6.14		
	MAMMA1000738	0.8	2.06	3.82	2.52	4.15	1.95		
	MAMMA1000744	1.12	2	5.52	3.27	3.97	4.67		
	MAMMA1000746	1	2.03	2.24	2.38	4.27	2.48		
40	MAMMA1000748	8.23	8.93	13.13	15.53	16.06	15.05	*	+
	MAMMA1000751	10.46	7.63	32.43	45.16	40.03	54.65	*	+
	MAMMA1000752	1.5	2.37	8.68	11.52	14.2	12.68	*	+
45	MAMMA1000757	1.89	2.48	5.54	7.9	8.53	10.86	*	+
	MAMMA1000760	3	2.99	6.77	4.65	8.01	7.12		
	MAMMA1000761	1.86	2.58	5.73	4.87	6.29	4.99		
	MAMMA1000775	1.37	1.83	4.43	3.23	4.29	2.64		
50	MAMMA1000776	2.37	2.36	6.3	6.57	7	6.12		
	MAMMA1000778	2.14	2.28	5.19	4.95	3.99	3.84		
	MAMMA1000781	1.33	1.33	3.06	2.82	2.86	1.23		
55	MAMMA1000782	1.94	2.36	3.88	2.93	3.11	3.01		
	MAMMA1000784	1.28	1.58	3.94	2.22	7.2	3.64		

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	MAMMA1000478	2.85	3.26	7.41	5.76	9.1	7.57		
	MAMMA1000483	4.16	3.16	8.3	8.09	6.5	8.63		
5	MAMMA1000490	1.65	2.61	3.68	2.66	4.96	2.14		
	MAMMA1000496	1.18	1.7	3.44	1.3	3.79	2.01		
	MAMMA1000500	0.68	1.79	3.22	1.41	3.2	2.86		
10	MAMMA1000501	3.04	3.89	7.86	13.71	15.02	12.51	**	+
	MAMMA1000503	0.84	2.08	2.21	3.52	3.52	2.27		
	MAMMA1000506	10.14	8.79	32.66	34.77	26.7	18.31		
	MAMMA1000510	3.24	3.5	4.59	10.97	10.76	13.61	**	+
15	MAMMA1000515	2.12	1.54	4.56	5.97	7.55	6.2	*	+
	MAMMA1000516	2.18	2.4	6.29	3.89	3.85	5.07		
	MAMMA1000522	1.04	1.47	4.39	2.4	3.68	1.85		
20	MAMMA1000524	2.04	2.09	3.53	3.82	6.18	3.96		
	MAMMA1000528	3.74	2.72	2.05	2.7	4.09	2.88		
	MAMMA1000534	0.91	2.35	2.11	1.86	2.91	2.04		
	MAMMA1000541	2.85	3.16	11.29	11.22	8.81	12.04		
25	MAMMA1000550	1.21	2.73	1.86	2.46	6.65	4.64		
	MAMMA1000556	1.78	1.32	4.25	2.66	3.37	1.4		
	MAMMA1000559	1.32	1.49	5.56	2.92	4.2	3.46		
30	MAMMA1000565	1.82	2.74	3.93	2.13	4.18	4.22		
	MAMMA1000567	0.99	2.16	3.77	2.3	4.07	3.64		
	MAMMA1000576	3.72	3.12	13.12	10.45	9.76	10.02		
	MAMMA1000582	2.07	2.7	5.64	4.13	4.81	6.31		
35	MAMMA1000583	1.16	2.33	2.45	2.19	4.47	2.93		
	MAMMA1000585	1.66	2.04	4.19	3.82	5.38	3.23		
	MAMMA1000587	1.64	1.51	3.73	3.12	5.07	3.49		
40	MAMMA1000591	0.96	1.34	3.11	1.45	3.74	1.72		
	MAMMA1000594	2.3	1.76	4.92	3.55	7.68	5.16		
	MAMMA1000597	4.42	3.09	9.64	9.46	9.63	10.35		
45	MAMMA1000605	2.84	3.94	11.44	18.34	15.85	17.89	*	+
	MAMMA1000612	1.91	2.15	5.22	3.85	4.33	4.95		
	MAMMA1000614	3.11	2.71	9.4	7.48	6.07	6.34		
	MAMMA1000616	1.66	1.79	2.44	2.1	5.09	3.45		
50	MAMMA1000621	1.39	1.67	3.36	3.15	6.31	7.02		
	MAMMA1000623	1.08	1.04	3.83	0.92	2.66	2.3		
	MAMMA1000625	7.39	6.32	23.76	19.68	25.39	29.8		
	MAMMA1000635	0.89	0.68	1.61	0.76	1.75	0.64		
55	MAMMA1000643	1.47	1.11	1.94	4.21	3.77	6.82	*	+
	MAMMA1000646	4.68	3.61	9.55	17.22	16.4	16.44	**	+

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	MAMMA1000331	1.08	1.65	3	3.33	6.01	3.21		
	MAMMA1000335	7.38	9.1	14.27	19.49	16.28	16.92	*	+
5	MAMMA1000339	0.33	0.39	-2.17	1.46	2.09	0.66		
	MAMMA1000340	1.43	1.33	4.12	4.37	2.72	2.15		
	MAMMA1000348	1.2	1.27	4.6	3.14	4.82	6.11		
	MAMMA1000356	1.93	2.21	4.93	3.08	5.42	6.29		
10	MAMMA1000358	2.93	3.97	5.02	7.42	7.47	4.9		
	MAMMA1000360	1.41	1.92	4.6	3.76	5.2	4.63		
	MAMMA1000361	2.2	3.45	8.9	8.94	10.67	8.93		
	MAMMA1000363	1.09	1.69	3.86	1.87	4.37	4.1		
15	MAMMA1000370	0.92	0.71	1.76	2.02	2.62	2.57	*	+
	MAMMA1000371	2.09	1.73	6.35	10.02	12.1	10.1	*	+
	MAMMA1000372	4.45	4.1	12.88	12.01	12.92	11.97		
20	MAMMA1000385	1.79	2.36	6.41	6.41	7.66	8.72		
	MAMMA1000388	1.93	3.02	6.03	4.7	4.53	5.06		
	MAMMA1000395	1.3	2.46	3.12	1.69	3.49	0.8		
25	MAMMA1000402	1.69	1.68	5.62	3.33	4.35	4.63		
	MAMMA1000403	1.7	2.36	5.05	5.45	5.81	3.96		
	MAMMA1000410	0.87	1.25	2.71	3.23	3.35	3.25	*	+
	MAMMA1000413	1.52	0.47	2.48	3.51	3.76	3.61	*	+
30	MAMMA1000414	1.08	1.53	3.03	2.94	4.91	1.81		
	MAMMA1000416	3.3	4.01	10.2	15.8	23.14	20.47	*	+
	MAMMA1000421	2.61	2.83	6.11	7.7	7.42	7.09	*	+
35	MAMMA1000422	2.83	2.53	7.46	9.18	6.64	12.05		
	MAMMA1000423	1.7	1.26	6	5.9	6.62	5.89		
	MAMMA1000424	0.88	1.7	3.17	1.91	2.38	1.07		
	MAMMA1000429	8.73	10.07	13.78	14.98	16.3	11.17		
40	MAMMA1000431	1.6	1.27	4.27	5.22	6.32	4.26		
	MAMMA1000432	1.05	2.33	2.85	2.63	2.82	1.41		
	MAMMA1000437	4.61	4.75	8.44	10.54	11.52	8.12		
45	MAMMA1000444	2.53	4.15	8.55	7.55	10.17	10.13		
	MAMMA1000446	1.19	2.07	3.87	2.03	3.63	2.49		
	MAMMA1000449	1.77	1.59	3.54	3.37	4.31	3.22		
	MAMMA1000457	4.44	4.82	7.12	7.2	6.88	6.22		
50	MAMMA1000458	1.27	2.22	4.83	2.52	4.03	1.94		
	MAMMA1000468	0.55	1.12	2.2	0.51	2.25	1.16		
	MAMMA1000472	1.15	2.3	4.42	4.77	6.36	5.79	*	+
55	MAMMA1000473	1.95	1.72	3.59	3.45	5.46	3.17		
	MAMMA1000477	3.86	3.29	5.67	8.71	9.92	7.97	**	+

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	MAMMA1000183	1.14	2.12	5.29	3.92	4.7	4.91		
	MAMMA1000191	3.25	3.34	14.68	12.56	14.7	18.23		
5	MAMMA1000192	5.76	8.53	-9.67	16.66	19.91	17.65	**	+
	MAMMA1000193	1.68	1.54	0.86	1.33	2.25	1.94		
	MAMMA1000198	1.88	1.99	5.53	4.44	6.49	5.57		
	MAMMA1000204	1.75	2.25	3.39	3.56	4.13	2.85		
10	MAMMA1000207	1	3.2	3.41	2.86	4.96	3.47		
	MAMMA1000214	1.76	2.08	3.68	2.84	4.42	4.74		
	MAMMA1000220	6.19	6.12	11.61	12.49	18.06	16.72	*	+
15	MAMMA1000221	0.57	1.04	1.68	1.14	4.51	0.87		
	MAMMA1000226	0.48	1.06	2.07	1.49	3.19	1.88		
	MAMMA1000227	0.93	1.23	1.6	2.73	3.67	3.46	**	+
	MAMMA1000230	1	1.23	1.77	2.38	3.04	2.94	**	+
20	MAMMA1000241	2.9	2.2	4.19	7.24	5.8	7.61	**	+
	MAMMA1000245	76.63	70.15	118.95	141.45	166.09	104.88		
	MAMMA1000248	6.79	4.17	13.48	13.18	13.44	18.8		
25	MAMMA1000251	1.68	1.72	4.7	5.55	5.39	5.29		
	MAMMA1000254	1.24	1.22	3.59	2.14	5.61	5.02		
	MAMMA1000257	5.39	2.62	25.06	32.2	43.78	35.79	*	+
	MAMMA1000262	15.48	9.75	18.2	40.81	33.23	34.89	**	+
30	MAMMA1000264	0.99	1.2	2.3	4.43	2.57	3.4	*	+
	MAMMA1000266	1.25	0.79	2.73	4.21	5.33	4.03	*	+
	MAMMA1000270	2.43	1.94	4.57	6.16	7.16	7.58	*	+
35	MAMMA1000271	6.01	3.26	8.54	8.94	6.17	8.1		
	MAMMA1000277	0.89	0.93	2.56	2.46	2.75	2.09		
	MAMMA1000278	1.84	2.01	4.29	2.18	5.06	3.51		
	MAMMA1000279	1.82	1.74	4.33	3.51	5.72	4.35		
40	MAMMA1000283	0.99	1.51	2.36	1.37	2.66	2.71		
	MAMMA1000284	2.65	2.51	8.31	6.28	8.49	8.01		
	MAMMA1000287	1.58	2.13	6.27	5.55	6.94	7.1		
45	MAMMA1000294	4.72	5.45	9.44	3.84	8.21	4.74		
	MAMMA1000298	0.87	1.36	2.51	1.55	3.1	0.95		
	MAMMA1000302	0.9	1.18	4.73	2.22	4.9	2.56		
	MAMMA1000303	0.92	1.62	2.63	4.16	4.06	3.22	*	+
50	MAMMA1000305	1.07	1.28	2.73	2.47	2.74	2.1		
	MAMMA1000307	2.29	3.03	9.61	15.85	14.04	14.38	*	+
	MAMMA1000309	0.57	1.61	3.69	4.6	3.65	4.61		
55	MAMMA1000312	3.55	4.99	8.08	6.19	5.18	6.55		
	MAMMA1000313	1.06	2.31	2.34	1.79	3.98	3.43		

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	LIVER1000185	2.42	2.55	5.16	4.37	4.72	4.23		
	LIVER1000187	0.96	1.55	4.84	6.64	4.17	3.5		
	LIVER1000190	3.77	3.48	5.95	4.71	6.24	4.47		
5	LIVER1000192	2.37	2.92	3.93	4.1	5.23	4.04		
	MAMMA1000009	1.39	2.55	5.12	3.62	4.72	3.46		
	MAMMA1000015	1.72	1.59	4.78	5.23	4.42	6.42		
10	MAMMA1000019	0.69	2.48	3.4	4.27	4.81	3.1		
	MAMMA1000020	2.79	2.35	5.63	6	7.86	5.75		
	MAMMA1000024	0.65	1.76	3.79	2.42	2.91	1.61		
	MAMMA1000025	1.92	2.56	6.92	4.96	6.72	5.6		
15	MAMMA1000043	1.06	2.36	6.43	6.93	8.22	6.6		
	MAMMA1000045	1.38	2.01	4.84	2.68	3.96	2.89		
	MAMMA1000046	1.74	2.44	3.18	2.88	4.5	2.37		
20	MAMMA1000055	8.51	8.71	9.57	9.38	10.74	9.36		
	MAMMA1000057	4.4	3.29	7.56	8.38	9.78	8.16		
	MAMMA1000060	26.78	24.33	45.25	48.69	33.84	48.6		
	MAMMA1000069	2.13	1.65	4.1	3.43	3.14	2.41		
25	MAMMA1000084	2.88	3	5.81	5.64	8.15	7.51		
	MAMMA1000085	2.75	3.74	7.02	6.45	5.82	6.97		
	MAMMA1000092	1.45	2.97	3.8	4.64	5.15	4.55	*	+
30	MAMMA1000096	4.45	4.96	9.29	8.15	9.11	6.09		
	MAMMA1000097	2.4	2.96	3.86	5.93	6.01	6.97	**	+
	MAMMA1000102	1.94	1.59	4.27	4.25	6.16	4.44		
	MAMMA1000103	1.52	1.65	5	2.39	4.56	2.83		
35	MAMMA1000106	1.25	2.15	5.1	2.3	4.48	3.27		
	MAMMA1000117	1.19	2.12	3.72	1.84	3.32	2.77		
	MAMMA1000118	1.03	2.06	3.08	3.38	3.21	4.56		
	MAMMA1000129	1.06	2.1	2.97	1.73	2.7	1.31		
40	MAMMA1000133	1.09	1.96	3.67	2.8	3.87	2.02		
	MAMMA1000134	1.23	2.08	4.28	2.27	4.61	1.93		
	MAMMA1000139	1.45	1.91	2.69	2.13	3.98	1.85		
45	MAMMA1000141	1.97	2.27	5.47	3.67	3.61	3.42		
	MAMMA1000143	1.66	1.1	2.55	3.83	3.52	1.74		
	MAMMA1000150	4.11	4.95	8.99	6.49	6.4	8.66		
	MAMMA1000155	1.87	2.71	4.35	5.46	5.69	6.41	*	+
50	MAMMA1000163	1.65	2.82	2.62	3.45	4.54	4.97	*	+
	MAMMA1000171	1.96	2.43	5.53	4.12	6.52	4.97		
	MAMMA1000173	3.5	5.27	10.33	17.47	17.44	18.26	**	+
55	MAMMA1000175	1.58	1.8	2.89	4.06	5.14	3.7	*	+

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	HEMBB1002712	8.55	1.32	2.38	2.92	4.06	1.4		
	IMR321000028	1.03	1.71	2.88	1.63	2.76	1.63		
5	IMR321000031	1.71	2.59	3.51	5.86	4.35	5.31	*	+
	IMR321000034	21.95	15.41	30.37	33.73	19.59	34.65		
	IMR321000039	5.81	7.11	14.41	14.72	15.71	13.99		
	IMR321000044	0.81	2.37	1.44	1.01	3.26	2.06		
10	IMR321000063	79.52	80.12	127.61	224.23	199.69	128.8		
	IMR321000085	21.02	18.07	26.38	30.28	48.13	47.89	*	+
	IMR321000089	1.51	1.42	3.86	3	6.7	5.84		
15	IMR321000091	4.79	2.91	6.5	8.35	11.38	8.55	*	+
	LIVER1000004	8.04	9.67	34.15	55.9	56.53	48.86	*	+
	LIVER1000008	1.13	1.36	3.06	1.68	4	2.17		
	LIVER1000011	3.03	5.9	26.65	37.8	54.37	45.77	*	+
20	LIVER1000022	2.75	3.66	7.75	9.39	9.82	9.17	*	+
	LIVER1000025	1.78	2.77	5.47	9.83	10.83	7.7	*	+
	LIVER1000030	1.05	0.96	2.12	2.04	2.56	1.23		
25	LIVER1000045	1.33	1.37	3.11	5.11	5.12	5.89	**	+
	LIVER1000046	1.01	1.53	3.86	4.14	7.82	5.34		
	LIVER1000072	1.61	1.26	5.23	12.42	9.54	12.21	**	+
	LIVER1000077	0.33	1.79	1.97	1.87	2.84	3.14		
30	LIVER1000080	1.53	3	5.81	5.96	4.24	5.41		
	LIVER1000086	6.38	7.69	47.4	69.84	79.87	70.57	*	+
	LIVER1000092	1.6	1.46	3.09	3.85	3.83	2.41		
	LIVER1000095	0.91	2.31	2.56	2.16	2.46	1.55		
35	LIVER1000097	1.26	0.74	2.49	2.18	2.84	2.25		
	LIVER1000098	0.43	1.37	2.57	2.76	3.95	2.29		
	LIVER1000100	3.3	2.82	5.82	4.99	7.44	3.74		
40	LIVER1000101	0.36	1.81	2.4	1.69	4.25	2.74		
	LIVER1000106	0.83	1.95	1.79	0.97	2.8	1.29		
	LIVER1000108	1.36	2.93	4.31	4.35	4.12	3.43		
	LIVER1000115	1.12	1.57	4.32	6.38	6.13	6.9	*	+
45	LIVER1000120	1.45	0.95	2.23	1.39	1.46	0.73		
	LIVER1000138	0.6	1.27	1.86	2.11	2.36	1.6		
	LIVER1000146	1.38	2.69	6.24	7.17	7.01	7.02		
	LIVER1000148	0.88	1.24	2.65	1.68	5.68	3.51		
50	LIVER1000157	30.11	26.71	67.66	123.41	85.61	124.96	*	+
	LIVER1000161	1.3	1.59	2.3	1.73	3.37	1.84		
	LIVER1000167	3.07	3.63	14.08	20.36	22.31	21.82	*	+
55	LIVER1000174	1.53	1.68	1.84	2.1	3.43	1.29		

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	HEMBB1002544	3.91	3.45	6.89	6.79	7.87	7.99		
	HEMBB1002545	0.92	2.76	2.83	3.21	4.15	4.29		
5	HEMBB1002550	1.32	1.69	1.86	2.99	4.68	2.42		
	HEMBB1002556	2.9	3.54	9.69	8.73	8.12	10.62		
	HEMBB1002571	17.25	14.03	19.8	21.91	16.59	24.61		
	HEMBB1002579	3.32	2.05	4.87	4.38	6.6	6.39		
10	HEMBB1002582	1.79	2.11	5.59	5.77	6.47	5.63		
	HEMBB1002584	2.82	1.94	6.09	3.94	4.67	3.62		
	HEMBB1002587	6.39	5.82	10.63	11.3	9.04	9.94		
	HEMBB1002590	1.6	3.07	7.46	5.86	7.3	5.84		
15	HEMBB1002596	1.5	2.01	3.17	5.59	4.94	4.21	*	+
	HEMBB1002600	1.55	2.72	3.81	5.02	7.18	3.93		
	HEMBB1002601	1.28	2.23	3.9	2.51	5.59	2.99		
20	HEMBB1002603	2.37	1.64	5.48	3.53	6.59	5.6		
	HEMBB1002607	1.48	1.15	4.34	2.59	4.26	2.99		
	HEMBB1002610	1.2	0.96	3.48	1.95	3.79	3.45		
	HEMBB1002613	0.96	2.41	4.31	3.98	5.39	3.72		
25	HEMBB1002614	3.18	3.34	5.35	3.87	6.08	9.76		
	HEMBB1002615	1.47	3.29	4.63	2.45	3.83	2.67		
	HEMBB1002617	0.67	3.09	2.88	2.1	3.34	3.4		
30	HEMBB1002623	2.31	3.63	4.36	3.96	6.28	5.91		
	HEMBB1002624	2.7	1.56	7.52	7.96	8.72	8.3		
	HEMBB1002631	1.65	2	4.28	2.14	3.72	1.74		
	HEMBB1002635	1.84	1.74	3.55	3.31	3.64	3.56		
35	HEMBB1002644	7.22	9.04	15.98	18.52	23.94	22.55	*	+
	HEMBB1002654	5.22	4.21	7.77	7.92	7.33	11.74		
	HEMBB1002661	1.93	2.16	3.96	1.99	4.13	5.33		
	HEMBB1002663	1.59	1.8	3.85	5.45	5.58	5.17	*	+
40	HEMBB1002664	1.28	2.4	4.43	5.05	9.22	6.76	*	+
	HEMBB1002677	1.88	1.83	1.86	1.81	4.79	2.34		
	HEMBB1002683	2.68	2.21	9.21	5.67	5.9	8.45		
45	HEMBB1002684	1.71	0.81	2.53	1.92	2.74	2.63		
	HEMBB1002686	1.23	1.39	2.88	1.45	3.37	1.64		
	HEMBB1002692	0.99	1.4	1.87	2.5	2.53	2.98	*	+
	HEMBB1002693	1.75	1.75	4.12	5.03	5.74	3.46		
50	HEMBB1002697	1.09	2.8	2.73	4.17	5.58	5.34	*	+
	HEMBB1002699	1.59	2.27	4.93	4.72	6.74	6.97		
	HEMBB1002702	1.63	1.5	2.54	1.76	3.25	3.33		
55	HEMBB1002705	4.2	2.84	6.79	8.83	8.26	7.92	*	+

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	HEMBS1002329	2.89	3	3.81	6.39	5.88	5.53	**	+
	HEMBS1002340	0.6	1.8	2.05	2.29	3.38	2.22		
5	HEMBS1002342	8.12	9.23	14.09	21.68	18.15	18.03	*	+
	HEMBS1002358	1.09	3.22	6.37	6.52	7.91	9.2		
	HEMBS1002359	1.09	2.55	4.29	4.66	5.4	3.54		
	HEMBS1002364	1.28	1.82	2.33	3.17	5.15	3.11		
10	HEMBS1002366	13.63	21.17	32.35	56.28	57.48	53.09	**	+
	HEMBS1002371	0.83	0.63	1.72	2.32	2.82	2.1	*	+
	HEMBS1002381	0.97	1.16	1.74	2.83	3.16	6.26		
15	HEMBS1002383	1.07	3.17	4.18	3.1	4.5	4.37		
	HEMBS1002387	0.98	2.36	2.68	2.09	3.68	2.96		
	HEMBS1002409	6.85	7.27	46.98	69.94	70.04	64.2	*	+
	HEMBS1002413	3.92	2.99	8.34	9.46	8.16	10.16		
20	HEMBS1002415	0.84	1.28	2.79	2.36	3.49	2.11		
	HEMBS1002424	1.04	1.17	1.63	2.89	3.04	2.66	**	+
	HEMBS1002425	1.12	1.69	5.86	6.46	10.1	5.26		
	HEMBS1002427	1.5	1.59	2.32	3.72	7.82	3.65		
25	HEMBS1002442	2.29	1.57	4.33	4.99	8.58	8.89	*	+
	HEMBS1002447	2.61	2.7	5.56	6.1	6.6	5.63		
	HEMBS1002453	2.5	2.48	6.56	6.31	7.55	5.25		
30	HEMBS1002457	1.54	2.08	4.77	3.69	4.61	4.8		
	HEMBS1002458	0.48	1.53	2.5	2.2	2.35	1.66		
	HEMBS1002463	1.36	1.84	6.55	6.24	6.11	8.87		
	HEMBS1002465	1.12	1.18	2.4	2.86	2.25	1.59		
35	HEMBS1002477	0.71	0.66	4.43	3	4.39	4.86		
	HEMBS1002479	22.08	21.58	27.54	16.12	19.41	17.27	*	-
	HEMBS1002489	0.86	3.02	3.9	5.73	5.51	7.68	*	+
40	HEMBS1002492	1.27	1.23	3.07	3.53	4.08	3.39	*	+
	HEMBS1002495	1.85	1.85	3.2	2.61	5.02	3.98		
	HEMBS1002502	0.94	2.52	2.81	1.77	4.83	3.27		
	HEMBS1002509	0.73	1.8	2.65	2.03	2.43	1.27		
45	HEMBS1002510	0.49	1.68	3.06	1.78	2.5	0.81		
	HEMBS1002520	1.46	2.47	5.44	6.62	7.57	8.61	*	+
	HEMBS1002522	0.82	1.88	4.42	2.31	6.8	2.07		
	HEMBS1002527	11.47	13.79	12.46	24.19	10.37	17.52		
50	HEMBS1002530	1.43	2.15	3.44	2.93	4.92	2.26		
	HEMBS1002531	0.46	1.32	2.04	1.23	2.99	0.35		
	HEMBS1002534	1.35	2.27	2.73	4.54	4.08	3.92	**	+
55	HEMBS1002536	6.58	5.93	46.38	45.93	63.71	42.88		

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	HEMBB1002092	2.51	2.24	4.48	5.34	3.65	6.27		
	HEMBB1002094	3.21	2.62	8.2	5.72	7.27	6.04		
5	HEMBB1002103	2.42	2.97	-3.51	3.74	5.58	4.47		
	HEMBB1002109	4.27	3.47	4.84	5.72	7.83	6.36	*	+
	HEMBB1002115	42.37	37.4	91.88	95.86	101.94	101.65		
	HEMBB1002120	0.89	1.22	2.91	0.86	2.94	2.41		
10	HEMBB1002121	0.75	1.56	1.63	1.5	4.66	2.53		
	HEMBB1002134	11.99	11.22	112.59	98.93	166.1	133.77		
	HEMBB1002136	1.29	1.65	2.9	2.59	3.26	2.58		
15	HEMBB1002138	10.48	9.64	20.72	18.78	23.06	19.4		
	HEMBB1002139	1.84	1.6	5.46	4.69	6	5.84		
	HEMBB1002141	1.53	0.83	3.44	1.48	4.64	2.54		
	HEMBB1002142	1.85	2	4.95	3.22	5.98	6.03		
20	HEMBB1002145	1.62	0.83	2.96	1.49	3.07	2.42		
	HEMBB1002152	1.27	1.19	3.15	2.32	6.36	3.41		
	HEMBB1002162	1.25	1.55	3.92	3.42	5.14	3.61		
	HEMBB1002173	4.18	1.09	5.58	2.77	4.48	3.84		
25	HEMBB1002189	2.78	1.95	6.14	7.01	8.25	5.93		
	HEMBB1002190	1.81	2.2	6.36	8.01	6.93	8.36	*	+
	HEMBB1002193	1.84	1.06	2.06	4.53	4.37	5.48	**	+
30	HEMBB1002217	3.82	2.26	6.02	3.61	6.06	3.23		
	HEMBB1002218	3.91	3.3	7.58	4.94	6.15	5.68		
	HEMBB1002228	2.28	2.9	6.17	6.68	7.94	6.97		
	HEMBB1002232	1.15	1.4	2.24	2.14	5.79	3.03		
35	HEMBB1002245	0.86	0.84	2.34	1.53	2.47	1.05		
	HEMBB1002247	1.72	0.59	2.44	1.38	2.24	1.03		
	HEMBB1002249	2.65	1.64	3.1	3.39	4.08	4.01	*	+
40	HEMBB1002254	1.35	1.35	3.83	2.64	3.27	2.86		
	HEMBB1002255	0.99	1.37	2.6	1.19	2.93	1.4		
	HEMBB1002266	1.33	0.83	2.07	0.73	1.99	0.54		
	HEMBB1002271	14.89	9.5	29.42	32.9	41.53	37.28	*	+
45	HEMBB1002280	1.62	0.78	1.6	1.55	2.93	1.12		
	HEMBB1002296	11.83	12.31	18.29	31.87	21.06	23.09	*	+
	HEMBB1002300	0.78	2.31	3.48	1.49	4.63	3.12		
	HEMBB1002302	1.17	2.17	3.26	2.49	4.76	3.04		
50	HEMBB1002306	1.83	1.96	4.28	2.91	4.34	2.6		
	HEMBB1002316	0.66	1.38	2.36	1.19	2.9	0.97		
	HEMBB1002326	0.93	1.68	4.52	4.35	4.41	3.9		
55	HEMBB1002327	0.99	0.99	2.66	1.46	2.95	2.06		

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	HEMBS1001952	1.41	2.05	4.72	2.31	4.73	3.1		
	HEMBS1001953	1.62	2.09	3.45	3.63	5.78	3.75		
5	HEMBS1001957	1.16	2.02	4.28	2.68	4.19	2.19		
	HEMBS1001959	2.05	3.31	4.07	4.22	5.75	3.68		
	HEMBS1001962	5.54	2.61	5.11	3.34	6.28	2.54		
	HEMBS1001967	2.59	2.46	5.62	6.11	8.49	6.15		
10	HEMBS1001973	2.25	2.4	6.14	7.62	10	7.28	*	+
	HEMBS1001978	2.08	1.71	6.29	7.2	7.57	5.83		
	HEMBS1001983	9.23	8.69	24.64	38.93	34.91	36.79	*	+
15	HEMBS1001987	1.78	2.34	3.64	1.66	4.75	2.38		
	HEMBS1001988	2.02	1.92	3.42	2.92	5.17	1.85		
	HEMBS1001990	7.65	7.72	9.18	12.44	11.53	15.42	*	+
	HEMBS1001996	1.54	1.47	3.89	1.61	3.51	1.22		
20	HEMBS1001997	1.46	2.25	6.1	4.2	5.98	4.23		
	HEMBS1001999	10.91	11.08	16.84	24.47	26.58	22.28	**	+
	HEMBS1002002	1.08	1.58	3.52	1.91	2.76	2.39		
25	HEMBS1002005	1.88	2.91	4.8	4.82	7.6	4.22		
	HEMBS1002009	2.32	2.48	3.03	2.24	6.23	2.7		
	HEMBS1002013	0.96	2.07	3.78	1.95	4.26	1.41		
	HEMBS1002015	3.95	4.25	9.47	5.82	8.92	6.73		
30	HEMBS1002024	45.16	34.47	111.32	113.31	106.76	120.55		
	HEMBS1002035	2.15	1.91	2.87	2.11	4.5	2.68		
	HEMBS1002039	1.18	2.29	5.1	3.28	5.9	2.98		
	HEMBS1002041	3.31	4.13	8.49	15.49	14.42	13.38	**	+
35	HEMBS1002042	3.97	4.66	9.49	8.09	10.63	9.94		
	HEMBS1002043	1.34	2.21	4.61	5.97	5.24	3.36		
	HEMBS1002044	0.4	1.19	2.68	1.25	4.19	1.92		
40	HEMBS1002045	2.83	2.5	10.03	6.34	7.63	4.56		
	HEMBS1002049	1.31	1.4	3.77	1.71	4.36	1.73		
	HEMBS1002050	1.62	1.61	4.5	3.31	4.53	2.94		
	HEMBS1002051	1.17	1.13	2.9	2.59	5.05	4.37		
45	HEMBS1002068	1.69	2.44	2.43	2.3	4.42	2.07		
	HEMBS1002069	3.39	3.94	7.83	6.86	7.55	5		
	HEMBS1002075	0.72	1.94	3.33	2.99	3.52	2.37		
	HEMBS1002079	1.2	1.8	1.89	1.22	2.84	1.3		
50	HEMBS1002080	1.74	1.85	4.78	1.55	6.02	2.41		
	HEMBS1002082	1.03	1.85	4.59	2.38	4.38	1.96		
	HEMBS1002084	25.86	22.68	51.44	33.52	35.54	38.37		
55	HEMBS1002088	13.92	15.78	22.14	29.46	37.25	35.66	**	+

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	HEMBS1001797	1.61	1.82	2.28	2.94	4.97	5.54	*	+
	HEMBS1001802	13.28	9.91	67.77	85.35	82.8	81.27	*	+
5	HEMBS1001812	2.32	2.58	-6.49	7.68	8.28	9.74	*	+
	HEMBS1001815	128.22	114.78	102.97	87.37	89.21	62.14	*	-
	HEMBS1001816	1.95	2.19	3.97	3.73	6.31	5.73		
	HEMBS1001831	0.69	0.98	1.54	0.72	3.28	1.16		
10	HEMBS1001834	16.15	9.68	133.91	102.49	173.48	141.04		
	HEMBS1001836	4.07	1.99	11.4	5.89	6.97	6.7		
	HEMBS1001839	0.89	0.86	1.43	1.08	1.95	1.21		
15	HEMBS1001841	80.32	59.68	120.73	35.74	39.04	23.13	*	-
	HEMBS1001844	5.26	4.72	9.73	10.15	10.68	8.11		
	HEMBS1001847	6.93	4.24	8.6	9.06	14.77	10.3		
	HEMBS1001848	25.33	21.68	40.92	49.73	73.94	62.17	*	+
20	HEMBS1001850	3.07	2.93	5.25	3.75	7.19	3.95		
	HEMBS1001859	13.4	8.82	20.85	39.61	27.01	40.75	*	+
	HEMBS1001863	1.7	3.65	7.66	6.89	8.88	7.5		
	HEMBS1001867	1.69	1.93	3.16	3.16	4.14	3		
25	HEMBS1001868	2.15	1.53	2.56	1.31	3.57	1.28		
	HEMBS1001869	1.5	2.3	4.23	4.62	8.15	3.58		
	HEMBS1001872	1.21	0.79	2.23	1.77	4.17	1.75		
30	HEMBS1001874	1.92	1.2	2.07	2.58	2.47	3.03	*	+
	HEMBS1001875	0.83	1.96	1.7	2.05	3.53	1.37		
	HEMBS1001880	2.68	2.17	6.79	4.41	8.91	8.35		
	HEMBS1001899	0.6	1.62	2.15	2.26	4.68	1.91		
35	HEMBS1001903	5.56	5.68	9.61	6.06	6.65	6.7		
	HEMBS1001905	2.04	2.82	5.49	4.98	6.07	7.14		
	HEMBS1001906	0.67	1.65	2.64	2.22	5.14	2.62		
	HEMBS1001908	1.82	1.63	5.26	3.31	6.56	3.59		
40	HEMBS1001910	1.92	1.16	2.64	3.2	4.41	4.61	*	+
	HEMBS1001911	2.06	1.11	4.2	5.73	4.81	4.33		
	HEMBS1001915	2.53	3.01	5.61	7.13	7.84	9.4	*	+
45	HEMBS1001921	2.19	2.21	7.12	6.56	8.47	8.25		
	HEMBS1001922	1.74	1.77	3.66	4.91	5.85	3.7		
	HEMBS1001925	1.48	2.24	4.57	3.07	4.73	3.39		
	HEMBS1001930	0.46	0.94	1.67	0.91	3.18	1.01		
50	HEMBS1001944	1.72	1.88	5.45	4.83	4.6	4.49		
	HEMBS1001945	2.15	1.05	3.58	2.25	3.97	2.12		
	HEMBS1001947	2.28	1.13	3.23	5.35	5.98	3.15		
55	HEMBS1001950	3.49	1.95	5.38	5.15	5.46	3.3		

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	HEMBS1001623	2.21	2.38	3.16	2.16	5.5	3.37		
	HEMBS1001625	3.73	3.04	4.33	2.93	4.81	3.79		
5	HEMBS1001630	1.31	2.36	3.54	1.84	4.3	1.23		
	HEMBS1001635	1.78	1.64	3.76	2.08	3.32	1.34		
	HEMBS1001637	1.76	1.14	3.98	2.09	4.4	3.58		
	HEMBS1001641	1.43	1.68	2.78	3.17	3.73	2		
10	HEMBS1001653	2.18	3.17	5.61	3.96	6.63	3.5		
	HEMBS1001665	1.08	2.17	2.04	2.5	4.43	0.88		
	HEMBS1001666	2.14	1.95	3.52	2.45	4.88	1.79		
15	HEMBS1001667	2.37	2.25	3.26	2.94	5.13	3.17		
	HEMBS1001668	3.19	2.11	5.15	2.45	6.42	2.69		
	HEMBS1001669	0.98	2.02	3.19	1.04	4.53	1.38		
	HEMBS1001670	4.02	4.82	6.88	10.7	9.71	8.65	*	+
20	HEMBS1001673	1.48	2.97	3.61	3.51	4.52	4.43		
	HEMBS1001675	1.83	3.27	4.65	4.68	5.78	4.88		
	HEMBS1001679	2.52	2.34	5.06	2.19	3.87	1.88		
25	HEMBS1001684	2.13	1.55	3.89	5.17	6.77	5.05	*	+
	HEMBS1001685	3.41	1.61	4.43	2.91	6.24	2.49		
	HEMBS1001695	1.9	2.22	4.43	1.38	3.88	2.12		
	HEMBS1001703	1.25	2.3	5.74	3.58	3.79	4.1		
30	HEMBS1001704	1.39	2.16	4.58	4.23	5.02	3.94		
	HEMBS1001706	2.76	2.6	3.58	5.6	6.26	4.87	**	+
	HEMBS1001707	1.35	2.01	2.87	2.25	3.67	2.8		
	HEMBS1001717	1.68	2.21	3.23	2.61	3.34	2.83		
35	HEMBS1001731	13.81	13.48	24.03	11.02	23.09	25		
	HEMBS1001734	3.47	3.35	7.62	6.88	9.22	4.18		
	HEMBS1001735	1.35	1.4	3.4	1.58	3.52	2.03		
40	HEMBS1001736	5.01	6.14	7.87	7.15	10.91	8.11		
	HEMBS1001747	0.92	1	3.23	1.87	3.67	2.82		
	HEMBS1001749	4.71	2.99	9.39	7.29	5.99	8.16		
	HEMBS1001753	3.79	3.3	5.5	7.4	8.97	9.3	**	+
45	HEMBS1001756	0.53	2.05	1.89	2.31	3.91	2.73		
	HEMBS1001757	1.08	1.8	2.64	3.04	4.86	4.54	*	+
	HEMBS1001760	1.32	0.98	3.74	1.49	3.56	2.13		
	HEMBS1001762	0.9	0.61	2.62	1.57	2.95	2.07		
50	HEMBS1001780	9.82	12.28	11.34	16.64	26.06	22.06	*	+
	HEMBS1001785	0.89	1.24	1.02	0.62	2.88	1.64		
	HEMBS1001788	3.22	1.26	5.17	5.76	6.13	5.3		
55	HEMBS1001793	5.6	4.73	18.12	22.08	20.38	22.86	*	+

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	HEMBS1001454	1.2	1.96	3.77	3.74	5.13	2.99		
	HEMBS1001458	4.72	6.48	12.41	7.88	9.04	6.89		
	HEMBS1001461	0.55	1.38	2.01	2.11	2.65	1.07		
5	HEMBS1001463	2.28	2.1	3.7	3.95	5.03	4.66	*	+
	HEMBS1001464	1.73	1.29	3.62	2.66	3.92	2.27		
	HEMBS1001466	1.15	1.84	2.75	1.88	3.77	2.73		
10	HEMBS1001482	1.76	2.21	4.36	2.94	5.36	3.97		
	HEMBS1001500	1.01	1.08	1.77	1.96	4.39	1.97		
	HEMBS1001505	3	3.32	5.87	9.35	14.06	11.17	*	+
	HEMBS1001521	2.06	2.43	5.4	5.15	5.78	7.14		
15	HEMBS1001527	2.63	4.74	11.16	8.69	9.66	9.6		
	HEMBS1001530	4.15	3.51	6.57	9.43	12.39	7.05		
	HEMBS1001531	1.11	1.34	4.62	2.99	5.16	4.17		
20	HEMBS1001532	0.63	1.86	2.77	1.86	4.41	2.15		
	HEMBS1001535	1.99	2.01	4.22	3.34	4.32	5.75		
	HEMBS1001536	2.18	2.65	6.37	4.62	6.87	5.45		
	HEMBS1001537	1.31	2.24	3.7	3.21	6.12	2.75		
25	HEMBS1001542	4.39	4.72	6.28	5.26	7.83	5.7		
	HEMBS1001543	7.84	3.58	8.49	8.13	7.08	5.38		
	HEMBS1001547	2.02	2.25	2.65	4.2	4.27	2.79	*	+
	HEMBS1001548	2.53	2.62	11.82	17.73	29.92	23.34	*	+
30	HEMBS1001551	0.89	1.7	4.47	2.93	5.96	2.65		
	HEMBS1001555	2.13	2.79	4.78	3.73	5.8	4.64		
	HEMBS1001562	1.9	2.64	4.27	2.23	3.46	3		
35	HEMBS1001564	132.08	140.08	310.28	333.18	233.12	279.03		
	HEMBS1001565	1.72	1.97	3.9	3.77	4.68	2.07		
	HEMBS1001569	0.79	0.8	3.04	2.49	3.68	1.44		
	HEMBS1001573	1.9	1.04	3.58	4.18	5.53	4.12	*	+
40	HEMBS1001585	1.5	1.96	10.91	3.75	7.14	4.87		
	HEMBS1001586	1.53	2.27	3.22	4.27	5.45	1.95		
	HEMBS1001588	1.33	2.9	5.74	4.72	6.32	5.06		
45	HEMBS1001595	2.68	3.33	6.92	3.78	4.84	4.7		
	HEMBS1001596	3.4	2.57	3.74	2.67	5.36	2.54		
	HEMBS1001599	1.45	1.57	3.21	3.07	3.47	2.06		
	HEMBS1001603	1.99	2.45	4.17	5.7	8.16	4.04		
50	HEMBS1001606	1.35	2.28	2.42	2	2.39	1.53		
	HEMBS1001612	4.31	3.07	9.25	8.81	8.09	8.9		
	HEMBS1001618	1.53	1.62	3.86	2.84	4.48	2.31		
55	HEMBS1001619	2.11	3.03	3.92	5.71	5.1	4.37	*	+

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	HEMBS1001289	4.64	5.82	12.2	6.93	9.11	6.88		
	HEMBS1001290	2.82	1.27	4.55	2.89	4.14	1.96		
5	HEMBS1001294	1.03	1.91	3.2	2.93	3.95	2.51		
	HEMBS1001299	7.06	7.64	12.49	16.26	14.41	17.87	*	+
	HEMBS1001302	2.16	2.34	2.41	1.75	3.39	1.64		
	HEMBS1001304	1.73	1.34	2.6	1.81	5.2	1.67		
10	HEMBS1001314	1.16	1.07	2.47	1.3	3.63	1.25		
	HEMBS1001315	1.25	1.62	1.46	0.87	4	0.9		
	HEMBS1001317	2.1	3.38	6.51	4.12	8.01	4.41		
15	HEMBS1001326	0.88	1.54	2.36	1.51	3	1.69		
	HEMBS1001331	2.11	2.79	2.81	3.78	6.14	4.55	*	+
	HEMBS1001335	1.39	0.9	1.44	1.4	2.81	1.92		
	HEMBS1001337	1.86	1.7	3.15	3.34	4.72	4.51	*	+
20	HEMBS1001339	4.17	3.87	5.91	5.83	7.91	5.25		
	HEMBS1001344	1.27	1.36	2.25	1.62	3.44	1.29		
	HEMBS1001346	2.17	2.32	7.45	5.89	6.23	5.7		
25	HEMBS1001348	0.68	1.37	4.05	1.38	3.68	2.82		
	HEMBS1001350	2.06	2.09	4.17	2.07	7.74	2.57		
	HEMBS1001356	1.4	1.9	2.33	1.74	5.54	2.29		
	HEMBS1001364	0.53	1.28	1.25	1.49	2.47	1.43		
30	HEMBS1001366	1.61	1.71	3.46	4.17	4.27	3.82	*	+
	HEMBS1001367	1.11	2.19	3.79	5.05	6.25	5.75	*	+
	HEMBS1001369	0.56	1.29	2.54	1.95	3.39	2.82		
	HEMBS1001380	3.13	3.56	6.65	5.43	7.36	7.75		
35	HEMBS1001381	8.45	6.07	9.53	10.2	14.39	11.86		
	HEMBS1001384	3.48	4.92	5.66	9.52	13.27	11.91	**	+
	HEMBS1001387	1.19	1.57	3.1	2.36	4.34	1.33		
40	HEMBS1001394	1.53	1.3	1.68	2.7	2.82	1.89	*	+
	HEMBS1001407	0.68	0.83	0.99	0.62	1.33	0.97		
	HEMBS1001410	1.35	1.04	1.78	2.44	2.63	1.74		
	HEMBS1001413	1.68	1.84	3.32	3.48	3.27	4.04		
45	HEMBS1001419	2.56	2.24	4.42	3.61	5.47	4.6		
	HEMBS1001421	2.29	1.66	2.18	1.31	3.16	0.95		
	HEMBS1001424	0.51	1.2	1.67	-0.1	1.58	0.41		
50	HEMBS1001426	2.04	1.51	3.7	2.66	5.67	4.21		
	HEMBS1001429	7.11	5.76	9.83	22.69	19.97	19.53	**	+
	HEMBS1001436	3.13	2.51	6.8	7.5	6.44	7.24		
	HEMBS1001443	5.61	6.48	20.67	20.46	27.07	22.15		
55	HEMBS1001449	2.02	2	4.92	4.26	6.35	4.27		

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	HEMBB1001126	1.85	1.88	3.63	2.69	4.96	2.9		
	HEMBB1001133	3.15	2.42	4.24	6.56	6.15	6.44	**	+
5	HEMBB1001137	1.97	2.2	4.4	3.28	6.56	5.42		
	HEMBB1001142	2.96	2.68	10.51	9.2	11.69	10.38		
	HEMBB1001145	3.25	3.56	7.39	6.11	7.7	6.59		
	HEMBB1001151	5.67	6.58	9.21	12.85	7.67	7.84		
10	HEMBB1001153	2	1.89	4.57	3.79	5.37	2.9		
	HEMBB1001158	6.96	6.74	12.17	12.04	9.95	11.16		
	HEMBB1001169	1.71	2.45	4.42	2.89	4.21	2.86		
15	HEMBB1001170	1	1.85	3.27	1.26	2.5	1.61		
	HEMBB1001175	1.43	1.54	5.16	3.39	6.36	4.14		
	HEMBB1001177	3.63	2.4	7.54	5.8	6.94	6.69		
	HEMBB1001182	2.6	3.69	4.89	3.89	6.54	3.87		
20	HEMBB1001192	3.3	3.09	16.1	15.27	20.06	15.75		
	HEMBB1001199	1.16	2.27	1.91	1.43	3.96	1.15		
	HEMBB1001200	1.86	1.66	3.14	2.43	5.38	2.29		
25	HEMBB1001208	2.02	2.04	4.56	2.96	6.19	2.74		
	HEMBB1001209	2.98	2.28	5.75	5.22	6.56	4.92		
	HEMBB1001210	5.14	4.28	7.8	11.05	6.08	10.33		
	HEMBB1001215	9.57	10.46	17.69	17.91	15.75	16.96		
30	HEMBB1001217	1.78	2.13	4.39	2.04	3.69	1.89		
	HEMBB1001218	4.28	3.37	5.47	4.52	5.98	4.6		
	HEMBB1001221	1.72	1.65	2.75	1.54	4.29	1.34		
	HEMBB1001224	2.2	2.46	3.81	2.98	6.08	3.7		
35	HEMBB1001230	1.51	2.09	4.36	2.55	4.95	2.22		
	HEMBB1001234	5.24	6.05	29.26	31	44.04	30.75		
	HEMBB1001235	12.72	10.54	21.49	13.27	8.47	10.71		
40	HEMBB1001237	11	10.54	21.03	32.1	26.16	36.86	*	+
	HEMBB1001242	4.82	5.68	8.63	6.92	6.97	4.51		
	HEMBB1001244	1.08	1.1	3.9	1.47	4.36	1.36		
	HEMBB1001249	1.26	1.63	2.99	1.84	5.52	1.98		
45	HEMBB1001253	1.53	1.92	5.68	1.96	4.89	2.39		
	HEMBB1001254	1.27	1.19	3.73	1.22	5.09	2.45		
	HEMBB1001266	2	4.32	4.75	4.49	6.09	4.63		
	HEMBB1001267	3.51	2.92	9.98	8.43	7.63	8.01		
50	HEMBB1001271	2.25	2.93	3.89	2.59	5.85	3.91		
	HEMBB1001282	2.27	2.68	3.9	2.77	4.86	2.99		
	HEMBB1001287	54.06	45.71	83.21	72.79	57.33	77.51		
55	HEMBB1001288	2.45	2.58	3.64	4.57	6.08	3.31		

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	HEM881000959	0.77	1.41	3.11	2.23	4.42	1.87		
	HEM881000973	1.21	1.12	2.6	2.07	4.47	3.2		
5	HEM881000975	0.9	1.28	-1.55	2.48	2.86	1.7	*	+
	HEM881000981	1.54	0.66	1.99	1.84	3.47	2.03		
	HEM881000985	1.67	1.82	2.71	3.59	3.99	3.24	*	+
	HEM881000991	0.99	1.35	2.83	1.46	3.75	2.43		
10	HEM881000996	4.89	3.12	6.78	6.75	6.64	11.18		
	HEM881001000	1.86	1.39	4.06	3.07	5.46	3.88		
	HEM881001004	1.15	1.32	2.57	2.56	4.55	2.42		
15	HEM881001008	1.48	1.79	2.75	2.44	4.6	2.88		
	HEM881001011	1.34	1.15	1.53	2.47	2.24	2.9	**	+
	HEM881001014	1.31	1.43	2.3	2.73	4.84	4.1	*	+
	HEM881001020	1.17	0.75	2.77	1.77	2.67	2.26		
20	HEM881001024	3.31	1.72	6.27	5.47	7.56	6.82		
	HEM881001026	5.14	4.03	5.16	5.46	7.67	5.44		
	HEM881001037	2	1.45	4.73	3.52	5.69	6.67		
25	HEM881001042	0.52	1.15	2.69	1.29	3.61	0.87		
	HEM881001046	1.18	1.28	2.16	1.67	3.82	0.96		
	HEM881001047	1.01	1.7	3.79	2.2	3.83	3.83		
	HEM881001048	2.5	2.34	7.02	4.34	11.02	6.93		
30	HEM881001051	1.44	2.62	3.23	3.95	6.26	3.9		
	HEM881001056	1.61	2.67	4.89	3.75	5.7	3.78		
	HEM881001058	1.3	1.92	4.72	2.64	6.92	2.63		
	HEM881001060	0.69	0.68	1.75	2.05	4.61	1.85		
35	HEM881001063	1.23	1.83	3.52	2.43	4.21	2.9		
	HEM881001068	1.84	3.62	3.59	3.46	7.14	5.2		
	HEM881001082	2.24	2.57	5.98	5.38	6.93	6.36		
40	HEM881001095	6.39	7.45	11.76	14.04	14.61	13.16	*	+
	HEM881001096	1.3	1.91	3.05	3.21	4.12	3.49		
	HEM881001101	7.41	8.19	9.74	19.33	13.1	16.69	*	+
	HEM881001102	1.04	1.47	4.57	3.6	6.07	4.46		
45	HEM881001104	1.66	1.89	3.98	3.1	4.87	4.25		
	HEM881001105	1.57	1.59	2.13	3.11	4.82	2.71		
	HEM881001112	9.44	8.91	73.3	100.88	136.14	131.28	*	+
50	HEM881001113	2.11	1.94	9.1	5.65	8.02	6.81		
	HEM881001114	1.88	2.27	5.18	4.16	7.06	4.82		
	HEM881001115	5.78	7.88	14.52	16.77	9.5	14.78		
	HEM881001117	1.7	1.52	2.92	1.85	2.79	1.79		
55	HEM881001119	1.69	1.57	4	2.29	3.74	2.28		

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	HEMBB1000794	1.46	1.8	2.85	2.97	3.84	2.06		
	HEMBB1000807	2.55	2.72	5.18	3.57	4	4.26		
5	HEMBB1000809	30.31	26.87	132.99	158.22	156.74	195.14	*	+
	HEMBB1000810	1.98	2.67	4.51	3.76	6.08	4.12		
	HEMBB1000821	1.98	1.93	2.98	2.05	4.45	1.79		
	HEMBB1000822	1.08	1.97	2.31	1.65	5.31	1.46		
10	HEMBB1000826	1.36	1.99	3.57	3.77	6.11	3.6		
	HEMBB1000827	2.48	2.89	5.83	2.67	5.05	2.99		
	HEMBB1000831	3.4	2.31	5.67	3.84	7.74	2.95		
15	HEMBB1000835	1.76	1.94	6.2	7.59	7.62	7.47	*	+
	HEMBB1000840	1.27	2.95	6.89	4.48	7.19	3.01		
	HEMBB1000848	2.08	3.45	5.63	5.39	6.45	5.3		
	HEMBB1000852	1.26	2.16	2.8	1.07	4.51	1.55		
20	HEMBB1000857	7.65	6.49	8.13	7.01	10.69	11.53		
	HEMBB1000858	3.7	3.13	7.3	7.07	9.38	7.31		
	HEMBB1000867	2.21	1.84	4.9	3.02	5.55	4.04		
	HEMBB1000870	1.64	2.37	4.56	2.84	5.31	3.63		
25	HEMBB1000876	1.48	2.86	3.91	4.54	3.22	3.93		
	HEMBB1000881	3.35	5.56	10.5	6.12	5.88	3.85		
	HEMBB1000883	1.02	2.68	2.2	3.03	3.32	2.58		
30	HEMBB1000887	16.9	14.54	43.41	67.39	61.26	59.84	*	+
	HEMBB1000888	1.03	1.67	2.39	1.63	3.92	1.86		
	HEMBB1000890	2.93	3.36	10.85	6.01	8.62	7.68		
	HEMBB1000893	3.28	2.54	5.46	4.5	6.14	5.57		
35	HEMBB1000900	1.27	1.53	2.98	2.06	2.54	1.58		
	HEMBB1000905	5.09	3.75	6.6	10.05	9.45	8.77	**	+
	HEMBB1000908	3.34	2.79	3.01	4.48	4.71	5.7	**	+
	HEMBB1000910	1.74	2.91	2.55	2.09	3.56	2.24		
40	HEMBB1000913	1.41	1.51	2.22	2.8	3.41	1.91		
	HEMBB1000915	32.08	25.6	50.05	48	58.92	51.07		
	HEMBB1000917	2.1	2.78	5.72	2.99	4.52	3.44		
45	HEMBB1000927	1.45	1.24	1.82	1.49	3.25	1.88		
	HEMBB1000932	0.66	2.06	2.74	1.81	3.41	1.61		
	HEMBB1000933	7.47	7.12	10.71	12.88	12.78	19.19		
	HEMBB1000936	1.44	1.96	2.87	3.75	6.44	3.55		
50	HEMBB1000939	7.86	7.14	9.02	15.98	15.3	18.25	**	+
	HEMBB1000941	1.53	1.86	3.17	3.99	4.46	3.52	*	+
	HEMBB1000947	3.53	3.34	4.61	4.67	6.8	5.72		
55	HEMBB1000954	1.08	1.82	2.54	1.62	4.01	3.08		

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	HEM881000611	0.94	1.16	2.08	1.15	2.79	1.66		
	HEM881000617	2.01	3.04	9.31	6.14	8.79	6.97		
5	HEM881000623	2.51	3.08	-4.64	5.58	5.83	3.9		
	HEM881000630	3.23	2.5	3.78	2.42	5.54	2.51		
	HEM881000631	8.91	10.69	18.75	22.52	23.76	22.55	*	+
	HEM881000632	6.77	8.77	20.85	27.2	18.4	23.31		
10	HEM881000636	9.52	15.91	22.42	25.26	21.65	19.96		
	HEM881000637	6.63	9.77	19.44	17.77	24.39	20.28		
	HEM881000638	1.44	1.41	3.23	3.6	5.29	3.34		
15	HEM881000642	3.47	2.31	7.58	7.65	9.33	9.93		
	HEM881000643	0.71	1.87	2.71	1.62	3.54	2.12		
	HEM881000649	2.25	2.22	6.45	4.94	7.61	5.72		
	HEM881000652	1.8	2.21	5.33	5.13	5.14	4.93		
20	HEM881000655	1.07	1.17	3.24	1.64	4.74	2.01		
	HEM881000665	0.52	1.08	2.23	1.69	2.92	2.01		
	HEM881000668	1.85	1.46	2.76	5.07	5.42	4.1	**	+
25	HEM881000671	2.36	2.01	6.77	7.03	7.81	6.94		
	HEM881000673	0.75	1.27	2.92	2.84	4.63	2.43		
	HEM881000679	3.26	2.84	5.59	4.42	7.19	5.76		
	HEM881000684	1.83	2.53	6.6	5.01	6.92	5.6		
30	HEM881000692	0.93	2	2.46	1.77	2.5	1.09		
	HEM881000693	0.96	1.29	2.47	1.6	2.79	1.34		
	HEM881000705	2.61	2.52	4.85	4.97	8.2	6.53		
35	HEM881000706	0.78	1.07	2.18	2.56	2.93	1.06		
	HEM881000709	3.53	2.92	8.39	8.16	7.99	10.26		
	HEM881000714	1.41	2.85	9.32	5.31	10.37	8.79		
	HEM881000725	1.61	2.22	4.35	3.04	6.22	4.72		
40	HEM881000726	1.88	2.34	8.76	5.63	7.1	4.83		
	HEM881000729	1.82	3.28	4.3	3.3	5.21	2.79		
	HEM881000738	1.94	2.6	5.55	3.99	5.53	6.15		
	HEM881000749	4.06	4.15	7.47	7.48	9.56	8.27		
45	HEM881000763	5.81	5.56	6.21	6.65	9.9	6.61		
	HEM881000770	2.76	2.06	8.8	7.73	9.62	8.83		
	HEM881000774	1.62	2.75	3.7	3.07	4.39	2.34		
50	HEM881000777	5.17	5.49	7.42	6.86	4.9	7.37		
	HEM881000781	3.7	4.19	6.89	7.64	5.28	6.83		
	HEM881000788	0.87	1.79	2.45	2.65	4.88	1.35		
	HEM881000789	1.91	2.22	3.01	3.1	6.55	1.86		
55	HEM881000790	1.97	2.15	4.48	4.59	4.21	2.53		

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	HEMBB1000430	38.77	36.24	61.06	51.76	34.69	50.02		
	HEMBB1000434	3.05	4.73	9.02	6.54	6.59	6.63		
5	HEMBB1000438	1.13	1.83	4.16	2.23	4	1.39		
	HEMBB1000441	2.26	3	7.35	5.44	8	4.78		
	HEMBB1000447	29.84	32.01	39.91	35.88	44.02	33.55		
	HEMBB1000449	1.3	1.31	3.72	1.51	3.04	1.54		
10	HEMBB1000453	8.61	8.04	13.39	14.23	18.78	13.74		
	HEMBB1000455	1.29	1.97	3.19	3.13	5.46	3.54		
	HEMBB1000472	2.3	2.28	4.22	4.07	4.35	3.52		
15	HEMBB1000480	1.9	3.59	7.03	5.71	6.63	5.87		
	HEMBB1000486	2.15	2.98	6.93	4.82	7.86	5.9		
	HEMBB1000487	1.21	1.79	4.48	2.66	4.8	2.57		
	HEMBB1000490	3.67	4.13	12.61	7.92	8.7	6.55		
20	HEMBB1000491	1.36	2.91	5	3.83	4.78	4.42		
	HEMBB1000492	3.02	4.04	6.84	5.63	6.94	5.34		
	HEMBB1000493	1.57	1.71	3.26	2.41	5.38	3.27		
	HEMBB1000510	1.32	1.71	4.94	4.4	5.61	4.21		
25	HEMBB1000516	5.64	7.71	36.22	16.62	18.58	17.09		
	HEMBB1000518	0.88	1.22	2.63	2.21	4.27	1.73		
	HEMBB1000523	1.32	2.78	7.41	3.33	7.74	4.1		
30	HEMBB1000530	2.83	2.51	9.72	6.06	7.81	6.64		
	HEMBB1000542	3.08	4.55	9.39	8.48	14.07	9.82		
	HEMBB1000550	4.84	2.87	4.77	10.48	5.74	5.33		
	HEMBB1000554	2.14	2.26	8.65	6.43	11.59	7.19		
35	HEMBB1000556	2.64	2.68	4.48	3.1	4.67	4.6		
	HEMBB1000564	1.81	1.4	5.87	4.26	5.34	6.18		
	HEMBB1000567	1.39	1.71	3.7	2.1	3.87	3.3		
40	HEMBB1000569	3.78	2.72	8.76	4.88	7.13	5.65		
	HEMBB1000573	3.48	3.44	11.31	7.33	9.55	7.56		
	HEMBB1000575	12.42	2.7	10.57	7.16	8.38	12.35		
	HEMBB1000579	2.12	3.75	4.84	3.01	8.61	4.04		
45	HEMBB1000585	0.83	1.19	3.19	2	4.1	3.35		
	HEMBB1000586	2.18	1.41	4.28	4.07	4.23	4.11		
	HEMBB1000589	2.98	1.58	3.62	3.38	3.75	3.69		
	HEMBB1000591	2.62	2.15	3.96	4.44	5.26	4.64	*	+
50	HEMBB1000592	2.05	1.2	3.18	3.87	3.39	2.15		
	HEMBB1000593	10.25	5.67	51.69	53.87	82.75	77.88	*	+
	HEMBB1000595	6.42	5.2	11.24	12.51	17.1	9.55		
55	HEMBB1000598	1.57	1.69	5.91	2.94	6.76	4.77		

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	HEMBS1000218	2.13	2.28	11.05	7.47	10.07	7.05		
	HEMBS1000226	2.63	3.26	6.02	4.02	5.86	3.41		
5	HEMBS1000230	1.39	1.83	-3.73	1.95	4.31	2.58		
	HEMBS1000240	6.04	8.26	10.01	3.97	2.7	1.75	*	-
	HEMBS1000244	1.39	1.64	3.51	2.51	2.63	1.68		
	HEMBS1000250	1.17	0.99	1.12	1.94	1.25	1.12		
10	HEMBS1000258	1.71	1.94	5.8	4.38	5.63	3.27		
	HEMBS1000264	2.49	3.12	11.01	8.64	8.34	8.1		
	HEMBS1000266	2.81	2.65	5.52	3.38	5.95	3.71		
15	HEMBS1000272	4.76	4.16	6.06	8.38	6.88	7.45	*	+
	HEMBS1000274	1.51	1.15	3.17	2.54	3.18	1.88		
	HEMBS1000276	1.12	1.84	4.72	3.1	4.01	2.43		
	HEMBS1000284	0.94	1.81	2.89	2.83	3.11	1.65		
20	HEMBS1000307	1.52	1.7	4.78	2.8	5.31	3.27		
	HEMBS1000309	1.43	2.73	3.07	3.09	3.56	2.19		
	HEMBS1000312	1.99	1.38	5.18	7.03	7.2	4.35		
25	HEMBS1000317	0.17	1.62	3.32	2.6	4.73	2.14		
	HEMBS1000318	1.11	2.69	3.85	2.28	4.46	1.68		
	HEMBS1000332	3.12	3.84	4.37	3.75	3.95	2.72		
	HEMBS1000335	0.77	2.35	4.66	6.16	4.66	3.44		
30	HEMBS1000336	0.99	1.11	3.59	2.09	3.52	2.29		
	HEMBS1000337	4.3	5.06	20.22	22.86	24.84	22.82		
	HEMBS1000338	2.11	1.92	5.86	7.13	8.92	4.71		
	HEMBS1000339	1.66	1.76	5.84	3.75	4.99	3.84		
35	HEMBS1000341	1.4	1.91	3.68	3.06	4.77	2.83		
	HEMBS1000343	2.51	3.15	6.96	7.24	8.68	7.46		
	HEMBS1000354	3.26	3.5	10.36	7	8.93	8.07		
40	HEMBS1000358	1.09	2.11	3.82	3.43	2.83	1.93		
	HEMBS1000369	1.93	2.33	3.87	5.96	6.54	2.89		
	HEMBS1000373	1.77	2.73	3.91	2.26	6.2	2.94		
	HEMBS1000374	3.27	4.06	9.34	9.58	13.36	6.95		
45	HEMBS1000376	2.71	3.92	12.28	10.03	6.99	7.71		
	HEMBS1000383	60.87	62.14	104.01	69.28	57.52	83.25		
	HEMBS1000391	1.8	2.66	4.57	4.89	6.18	4.29		
	HEMBS1000399	2.51	3.79	3.69	3.93	5.72	3.71		
50	HEMBS1000402	1.61	2.06	3.33	2.67	5.3	1.72		
	HEMBS1000404	1.34	1.15	5.18	2.56	5.29	1.81		
	HEMBS1000407	2.2	3.36	6.76	5.57	5.75	4.66		
55	HEMBS1000420	1.93	1.46	3.86	4.33	4.76	4.54	*	+

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	HEMBB1000030	3.12	3.53	6.58	6.62	7.77	6		
	HEMBB1000036	5.3	4.76	5.04	6.95	8.19	5.93	*	+
5	HEMBB1000037	4.43	3.64	4.73	4.63	8.38	5.32		
	HEMBB1000039	1.17	0.96	3.98	2.61	4.11	2.95		
	HEMBB1000044	1.22	2.35	4.26	5.28	5.58	5.36	*	+
	HEMBB1000048	3.2	1.7	3.48	3.99	5.4	3.96		
10	HEMBB1000050	2.32	1.55	3.33	2.97	3.98	2.85		
	HEMBB1000054	2.03	2.08	7.07	4.49	5.09	3.98		
	HEMBB1000055	42.59	36.75	92.41	100.33	86.52	89.35		
	HEMBB1000059	2.5	2.65	11.34	10.96	11.52	14.73		
15	HEMBB1000072	6.84	7.77	58.85	73.22	97.61	76.22	*	+
	HEMBB1000081	2.85	3.56	10.79	5.69	6.19	7.23		
	HEMBB1000083	1.13	1.95	5.38	4.88	5.89	5.33		
20	HEMBB1000089	1.14	2.53	4.54	6.03	6.73	5.43	*	+
	HEMBB1000094	4.12	4.3	8.4	4.24	5.04	6.62		
	HEMBB1000097	2.48	1.71	7.91	4.75	4.55	4.3		
	HEMBB1000099	2.69	2.07	6.27	5.18	6.75	5.64		
25	HEMBB1000103	7.19	5.28	18.55	13.99	19.26	16.16		
	HEMBB1000106	3.91	3.75	8.15	4.24	6.4	5.9		
	HEMBB1000113	1.25	1.54	3.33	1.39	4.14	2.31		
30	HEMBB1000119	2.19	2.17	5.66	3.34	6.12	4.05		
	HEMBB1000133	21.01	22.21	30.57	43.5	66.13	60.69	*	+
	HEMBB1000134	4.92	2.95	8.69	13.39	9.79	9.8		
	HEMBB1000136	7.14	8.81	29.63	23.11	26.28	28.23		
35	HEMBB1000141	1.98	2.85	6.18	4.95	5.75	6.16		
	HEMBB1000144	2.05	2.59	4.85	3.09	5.36	1.38		
	HEMBB1000147	3.77	2.08	4.51	4.55	7.39	3.07		
40	HEMBB1000152	0.79	1.45	3.42	5.13	5.15	3.45		
	HEMBB1000154	0.98	1.11	3.43	2.59	2.89	2.47		
	HEMBB1000155	0.88	0.54	3.15	3.33	4.15	1.92		
	HEMBB1000173	3.35	3.72	12.14	10.1	10.89	7.51		
45	HEMBB1000175	1.85	1.32	3.39	6.06	4.09	3.86		
	HEMBB1000176	1.48	4.03	6.12	3.43	9.75	5.03		
	HEMBB1000198	0.88	1.72	3.64	2.6	3.59	3.22		
	HEMBB1000208	1.12	1.52	3.04	1.74	3.23	2.69		
50	HEMBB1000209	1.62	1.54	3.76	3.32	3.94	3.28		
	HEMBB1000212	1.88	1.03	3.26	4.37	3.93	1.98		
	HEMBB1000215	1.61	2.13	3.8	4.67	5.49	5.32	*	+
55	HEMBB1000217	5.67	4.97	11.23	15.21	18.81	11.78	*	+

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	HEMBA1007186	1.71	2.76	4.49	4.95	5.47	3.86		
	HEMBA1007194	4.81	3.43	5.58	7.83	9.34	8.67	**	+
5	HEMBA1007200	1.18	2.33	-- 2.9	3.25	4.6	1.66		
	HEMBA1007203	1.54	3.5	5.38	6.03	6.87	5.05		
	HEMBA1007206	1.92	2.46	5.72	7.07	7.91	5.94		
	HEMBA1007224	5.4	6.5	9.06	9.23	5.85	8.06		
10	HEMBA1007226	7.19	8.07	40.61	59.31	70.51	62.19	*	+
	HEMBA1007240	10	10.96	13.45	15.35	7.71	11		
	HEMBA1007241	3.59	2.88	4.56	4.61	6.81	3.63		
15	HEMBA1007242	2.52	2.86	5.01	6.29	6.87	4.23		
	HEMBA1007243	10.23	10.91	69.57	70.17	95.69	82.75		
	HEMBA1007251	1.32	1.8	4.14	3.02	3.67	2.01		
	HEMBA1007256	1.39	1.91	3.36	3.93	5.74	3.44		
20	HEMBA1007267	3.19	3.71	8.75	8.73	9.18	8.1		
	HEMBA1007273	0.98	2.66	3.84	3.56	5.82	2.25		
	HEMBA1007279	1.55	2.25	3.52	2.95	4.35	2.02		
25	HEMBA1007281	1.73	1.54	2.12	2.95	4.43	1.01		
	HEMBA1007283	2.45	3.15	6.78	6.37	6.58	5.96		
	HEMBA1007288	2.12	2.77	5.54	4.35	6.74	5.48		
	HEMBA1007291	1.59	1.8	4.29	2.14	4.4	0.98		
30	HEMBA1007299	20.39	22.25	39.67	40.95	47.97	40.26		
	HEMBA1007300	2.08	2.75	3.59	4.17	4.45	4.07	*	+
	HEMBA1007301	1.97	2.82	3.15	3.73	3.99	3.44	*	+
	HEMBA1007319	2.84	3.61	6.73	5.21	6.12	3.32		
35	HEMBA1007320	1.29	1.22	3.12	4.19	3.45	2.42		
	HEMBA1007322	19.97	17.81	27.74	45.24	39.42	37.31	**	+
	HEMBA1007323	4.54	6.69	11.47	6	6.36	6.21		
40	HEMBA1007326	4.58	3.85	13.34	8.29	8.07	9		
	HEMBA1007327	3.37	3.98	8.91	6.14	9.31	8.98		
	HEMBA1007332	3.12	3.47	5.42	5.27	7.56	5.33		
	HEMBA1007341	1.4	2.51	3.24	2.93	3.38	3.36		
45	HEMBA1007342	1.06	2.05	3.02	1.52	3.23	1.98		
	HEMBA1007347	3.39	3.24	6.34	4.55	7.34	6.32		
	HEMBA1007353	2.43	2.22	4.99	2.68	6.9	3.01		
50	HEMBA1000005	1.57	2.54	5.35	2.68	4.65	3.64		
	HEMBA1000008	2.19	2.53	5.99	3.51	6.31	3.71		
	HEMBA1000018	2.21	2.16	7.13	9.9	9.79	7.24		
	HEMBA1000024	3.71	2.15	5.4	5.13	5.77	6.39		
55	HEMBA1000025	2.11	2.09	3.55	1.68	5.13	2.62		

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	HEMBA1007002	7.02	4.12	31.4	26.92	38.45	42.11		
	HEMBA1007013	1.02	0.94	3.04	1.44	4.39	2.27		
5	HEMBA1007016	2.02	1.43	-5.06	3.27	5.97	5.28		
	HEMBA1007017	0.69	1.24	2.55	1.42	3.33	1.83		
	HEMBA1007018	4.02	4.52	6.54	7.65	6.1	7.07		
	HEMBA1007044	8.13	8.41	17.4	15.48	11.97	12.27		
10	HEMBA1007045	1.64	2.15	4.42	2.61	5.08	3.47		
	HEMBA1007051	2.26	2.56	4.71	3.42	4.42	3.28		
	HEMBA1007052	2.23	1.25	3.47	2.37	4.62	1.83		
	HEMBA1007053	1.83	3.14	4.03	2.64	4	2.5		
15	HEMBA1007057	0.92	2.56	3.21	3.52	4.03	3.41		
	HEMBA1007062	0.91	0.82	2.73	2.34	2.87	1.82		
	HEMBA1007063	3.87	2.7	8.87	8.56	8.02	7.58		
20	HEMBA1007066	1.72	2.03	3.62	2.2	4.39	2.71		
	HEMBA1007069	1.36	2.29	3.87	3.84	3.48	3.66		
	HEMBA1007073	1.93	2.16	6.12	2.96	9.57	3.6		
	HEMBA1007076	1.48	2.6	5.56	4.66	7.6	3.4		
25	HEMBA1007078	6.5	6.83	18.61	26.96	27.47	23.78	*	+
	HEMBA1007080	7.6	9.46	38.27	61.02	68.15	51.34	*	+
	HEMBA1007084	1.28	1.42	4.76	3.68	6.27	4.76		
30	HEMBA1007085	3.28	2.89	8.21	5.76	7.21	5.42		
	HEMBA1007087	2.6	2.88	6.31	3.96	6.92	5.63		
	HEMBA1007089	26.17	28.1	43.8	46.11	34.41	29.42		
	HEMBA1007095	75.81	62.79	111.43	134.53	48.47	121.4		
35	HEMBA1007101	2.78	3.27	8	26.73	21.52	19.57	**	+
	HEMBA1007104	1.87	1.92	3.52	2.46	4.53	2.57		
	HEMBA1007106	4.77	4.8	9.03	16.42	12.5	9.49		
	HEMBA1007112	3.01	3.07	5.16	6.39	6	4.77		
40	HEMBA1007113	1.53	2.29	9.04	6.03	6.1	5.97		
	HEMBA1007121	13.76	14	92.08	116.14	111.53	129.12	*	+
	HEMBA1007129	1.54	2.44	2.87	2.66	4.37	1.89		
45	HEMBA1007147	1.68	2.88	4.4	3.96	4.76	4.06		
	HEMBA1007149	5.3	7.24	8.38	10.48	6.82	9.73		
	HEMBA1007151	0.85	1.87	3.38	3.32	3.88	2.54		
	HEMBA1007172	1.26	1.91	4.13	2.96	4.81	3.51		
50	HEMBA1007174	1.4	1.43	2.75	3.96	3.65	2.5		
	HEMBA1007176	2.58	3.95	11.7	6.7	6.78	4.52		
	HEMBA1007178	4.77	4.71	9.32	10.94	13.03	8.12		
55	HEMBA1007185	9.38	10.32	9.59	19.5	7.83	15.16		

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	HEMBA1006843	19.27	17.89	35.47	55.34	39.67	68.17	*	+
	HEMBA1006849	5.64	4.37	11.23	10.07	11.23	10.83		
5	HEMBA1006850	31.45	33.76	-60.2	44.62	53.25	45.59		
	HEMBA1006861	12.19	11.3	24.61	22.49	16.43	17.99		
	HEMBA1006865	5.42	8.35	31	33.77	33.36	34.49		
	HEMBA1006867	4.32	5.03	6.41	6.27	7.76	6.15		
10	HEMBA1006873	3.14	3.59	8.87	6.5	9.26	7.75		
	HEMBA1006877	2.52	4.03	5.87	4.03	6.26	3.68		
	HEMBA1006878	2.52	2.36	6.79	3.82	7.86	3.51		
15	HEMBA1006879	6.19	5.68	8.83	11.06	15.86	17.33	*	+
	HEMBA1006884	10.11	3.17	6.59	6.15	8.42	8.78		
	HEMBA1006885	7.02	6.82	14.16	20.86	19.11	21.73	*	+
	HEMBA1006886	20.38	17.25	26.55	29.45	29.15	40.93		
20	HEMBA1006889	2.61	4.02	4.85	4.48	5.99	6.09		
	HEMBA1006896	19.11	24.76	31.7	33.67	39.11	40.41	*	+
	HEMBA1006900	6.19	5.93	20.76	16.81	23.73	18.45		
25	HEMBA1006902	1.43	2.45	3.86	4.03	6	3.98		
	HEMBA1006912	1.24	1.74	6.86	4.12	5.8	5.3		
	HEMBA1006914	6.64	6.11	18.27	14.81	18.62	15.03		
	HEMBA1006916	3.11	2.71	5.78	10.29	7.48	9.36	*	+
30	HEMBA1006921	3.03	3.5	9.63	9.77	11.26	13.59		
	HEMBA1006926	2.65	2.61	5.68	5.01	6.53	6.98		
	HEMBA1006927	3.06	2.2	5.17	3.57	5.26	5.89		
	HEMBA1006929	2.94	2.69	4.02	4.31	6.36	5.25	*	+
35	HEMBA1006936	3.72	3.21	6.51	4.67	6.25	5.45		
	HEMBA1006938	1.21	2.11	6.57	2.37	3.76	3.44		
	HEMBA1006941	9.52	8.15	12	19.26	28.62	23.74	**	+
40	HEMBA1006942	5.2	2.63	6.65	10.7	10.65	11.4	**	+
	HEMBA1006945	10.07	5.91	16.81	23.73	17.09	19.91		
	HEMBA1006949	1.6	1.43	3.88	2.48	5.34	2.81		
	HEMBA1006952	1.16	1.66	2.98	3.02	5.04	2.22		
45	HEMBA1006960	2.53	2.78	7.66	5.9	8.28	8.68		
	HEMBA1006973	1.74	2.27	5.91	4.7	7.84	5.54		
	HEMBA1006974	2.49	3.44	6.76	6.09	11.01	8.14		
	HEMBA1006976	1.39	1.5	4.12	3.18	4.96	4.36		
50	HEMBA1006989	1.85	1.66	6.51	2.05	3.01	1.81		
	HEMBA1006993	2.71	2.39	6.49	6.11	7.69	8.79		
	HEMBA1006996	0.74	1.15	2.98	2.52	3.13	3.58		
55	HEMBA1007001	1.91	2.47	5.12	3.98	6.37	4.76		

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	HEMBA1006666	1.83	1.25	3.23	3.24	4.14	2.45		
	HEMBA1006671	8.69	6.74	11.66	17.25	16.13	15.21	**	+
5	HEMBA1006674	1.64	1.99	7.42	6.71	9.73	7.34		
	HEMBA1006676	1.46	2.36	5.19	4.28	6.75	3.19		
	HEMBA1006682	2.17	1.64	3.43	2.2	4.98	1.88		
	HEMBA1006688	1.48	2.46	4.74	3.31	4.38	3.54		
10	HEMBA1006695	1.58	2.41	4.85	3.54	5.79	2.46		
	HEMBA1006696	2.84	3.93	6.29	5.95	6.74	5.82		
	HEMBA1006702	3.31	1.83	13.28	4.13	5.58	4.87		
	HEMBA1006707	2.89	2.62	5.9	5.04	7.54	5.5		
15	HEMBA1006708	2.21	1.52	5.71	4.42	4.57	2.63		
	HEMBA1006709	1.64	1.97	4.6	4.29	5	4.74		
	HEMBA1006717	1.58	2.28	3.58	2.5	4.93	2.28		
20	HEMBA1006724	2.68	3.42	4.55	4.45	4.5	5.47		
	HEMBA1006731	1.83	2.95	3.95	4.12	5.51	3.1		
	HEMBA1006737	1.82	3.5	6.59	3.89	5.09	4.45		
	HEMBA1006742	1.78	2.44	4.16	3.32	4.14	3.61		
25	HEMBA1006743	4	4.02	11.48	14.16	17.25	11.88		
	HEMBA1006744	1.84	1.79	7.74	6.6	8.29	5.72		
	HEMBA1006749	1.14	1.27	3.72	1.88	3.8	1.71		
30	HEMBA1006752	16.53	16.28	26.81	35.31	18.85	33.99		
	HEMBA1006754	1.44	2.6	3.63	5.55	4.33	2.49		
	HEMBA1006758	1.38	2.83	4.25	6.89	4.68	4.02		
	HEMBA1006767	3	4.14	7.88	6.81	7.89	5.51		
35	HEMBA1006770	5.05	2.61	7.12	7.89	8.1	5.05		
	HEMBA1006779	4.44	4.1	10.99	9.57	10.28	8.42		
	HEMBA1006780	3.28	3.19	10.27	7.6	8.33	8.38		
	HEMBA1006789	2.83	1.87	11.34	4.55	5.42	5.51		
40	HEMBA1006795	2.13	2.45	7.58	4.56	7.87	3.29		
	HEMBA1006796	4.31	3.15	5.37	5.85	7.06	4.72		
	HEMBA1006805	2.72	2.73	5.9	12.77	17.38	12.96	**	+
45	HEMBA1006807	30.32	28.07	75.38	54.22	54.72	67.82		
	HEMBA1006813	0.93	1.73	2.81	1.93	4.17	0.97		
	HEMBA1006819	3.73	4.53	8.5	7.02	8.09	5.29		
	HEMBA1006821	1.56	2.37	6.09	5.05	4.24	4.36		
50	HEMBA1006824	2.13	3.13	7.39	5.69	6.04	5.75		
	HEMBA1006832	19.84	18.63	56.97	59.26	61.56	53.28		
	HEMBA1006834	13.23	12.47	20.38	29.88	28.37	21.62	*	+
55	HEMBA1006835	1.11	1.49	3.88	5.08	7.25	4.6	*	+

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	HEMBA1006492	5.34	7.71	10.69	15.16	24.33	15.92	*	+
	HEMBA1006494	1.27	1.18	2.67	2.44	3.88	1.86		
5	HEMBA1006497	1.67	2.33	-4.75	4.3	3.79	4.49		
	HEMBA1006501	7.61	7.52	62.05	58.03	78.37	58.59		
	HEMBA1006502	4.73	3.55	15.72	18.66	22.55	21.66	*	+
	HEMBA1006507	8.7	6.8	51	49.69	71.2	48.41		
10	HEMBA1006517	1.51	1.99	5.07	3.43	5.9	4.64		
	HEMBA1006521	1.79	1.8	4	2.41	4.55	3.02		
	HEMBA1006529	4.77	3.74	3.86	5.97	5.83	2.78		
15	HEMBA1006530	1.8	1.39	2.06	1.62	3.53	2.42		
	HEMBA1006535	1.66	1.43	2.01	2.66	2.81	2.43	**	+
	HEMBA1006536	0.59	2.22	3.96	3.04	3.23	2.33		
	HEMBA1006540	1.61	1.68	3.33	3.05	4.1	3.56		
20	HEMBA1006544	1.39	1.63	8	3.54	5.85	4.35		
	HEMBA1006546	2.06	2.56	6.98	4.25	5.77	4.51		
	HEMBA1006549	1.74	2.13	5.93	4.57	4.63	4.61		
25	HEMBA1006559	2.55	1.45	4.63	2.99	5.76	3.32		
	HEMBA1006562	0.74	1.32	4.07	2.39	5.24	2.72		
	HEMBA1006566	0.67	1.28	0.97	1.34	1.69	0.99		
	HEMBA1006569	2.33	1.36	3.97	3.25	3.89	4.02		
30	HEMBA1006572	1.02	2.38	2.94	2.68	3.92	2.01		
	HEMBA1006579	20.44	16.82	51.93	59.14	66.28	72.54	*	+
	HEMBA1006583	3.17	2.85	5.59	5.83	5.94	6.45		
35	HEMBA1006595	1.82	1.63	4.29	2.83	5.22	3.68		
	HEMBA1006597	1.65	2.1	5.34	3.81	6.75	3.77		
	HEMBA1006606	1.75	2.31	4.96	3.72	5.71	2.8		
	HEMBA1006612	2.63	3.54	7.11	7.32	5.31	7.07		
40	HEMBA1006617	1.93	2.58	6.24	4.62	4.6	5.41		
	HEMBA1006624	6.37	8.61	16.93	12.84	15.78	10.91		
	HEMBA1006631	3.24	3.13	8.01	8.11	11.54	6.97		
	HEMBA1006635	1.7	2.57	5.7	4.27	6.27	3.8		
45	HEMBA1006639	1.12	1.98	4.52	4.14	5.3	3.06		
	HEMBA1006643	1.88	1.23	1.85	2.12	3.81	2.17		
	HEMBA1006648	7.69	6.84	16.92	17.63	23.57	23.73	*	+
50	HEMBA1006652	5.96	7.86	16.11	16.41	16.83	17.34		
	HEMBA1006653	2.21	3.66	7.23	6.92	5.18	5.17		
	HEMBA1006658	5.04	6.58	12.68	14.07	13.59	12.01		
	HEMBA1006659	6.8	10.14	48.92	66.47	78.17	67.44	*	+
55	HEMBA1006665	1.44	0.89	3.32	2.77	2.91	1.82		

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	HEMBA1006347	2.04	1.83	4.2	2.94	2.76	1.68		
	HEMBA1006349	2.47	2.79	6.73	3.62	5.65	3.5		
5	HEMBA1006352	1.65	1.65	3.27	2.63	5.64	1.32		
	HEMBA1006357	4.99	4.26	8.6	8.36	7.8	7.74		
	HEMBA1006358	1.67	2.57	4.95	4.7	5.11	5.63		
	HEMBA1006359	1.56	2.17	4.02	3.67	5.48	3.3		
10	HEMBA1006360	2.53	2.12	5.1	9.73	10.02	8.61	**	+
	HEMBA1006364	1.71	2.76	4.82	4.02	5.55	3.51		
	HEMBA1006377	5.8	8.03	13.25	20.48	16.19	15.31	*	+
15	HEMBA1006380	1.31	1.57	8.75	6.78	7.01	6.9		
	HEMBA1006381	2.38	3.07	12.65	6.18	7.97	6.66		
	HEMBA1006385	3.21	3.33	8.65	5.41	8.17	5.32		
	HEMBA1006390	9.49	7.85	14.66	22.01	20.52	22.7	**	+
20	HEMBA1006391	6.58	6.85	6.73	12.83	10.12	13.15	**	+
	HEMBA1006398	1.32	1.67	4.19	2.57	4.12	1.69		
	HEMBA1006405	23.81	23.5	38.82	23.85	22.13	24.92		
25	HEMBA1006410	8.26	4.16	6.14	5.74	11.61	6.06		
	HEMBA1006416	2.14	2.62	5.93	5.3	6.98	4.23		
	HEMBA1006418	5.06	5.49	11.76	8.17	8.09	5.92		
	HEMBA1006419	2.67	3.93	8	6.89	7.77	5.2		
30	HEMBA1006421	2.03	3.28	3.09	3.44	4.19	2.27		
	HEMBA1006424	1.48	1.92	3.59	1.94	5.42	1.84		
	HEMBA1006426	3.03	3.99	7.91	7.23	7.87	5.51		
	HEMBA1006430	2.31	2.64	6.29	5.89	7.43	5.08		
35	HEMBA1006438	2.06	2.22	6.35	4.37	6.53	2.92		
	HEMBA1006445	1.98	2.68	5.72	6.11	5.6	4.16		
	HEMBA1006446	1.32	2.61	5.59	2	6.51	2.43		
40	HEMBA1006456	3.51	5.07	8.64	14.9	21.76	15.91	*	+
	HEMBA1006461	1.54	2.18	5.35	4.35	5.49	4.07		
	HEMBA1006467	1.52	1.78	3.61	2.82	7.24	3.2		
	HEMBA1006470	4.06	4.03	22.46	18.72	29.54	19.52		
45	HEMBA1006471	1.58	1.6	6.6	6.45	6.78	6.26		
	HEMBA1006474	7.35	6.37	43.12	55.87	62.01	52.31	*	+
	HEMBA1006476	9.48	10.05	66.66	94.3	119.21	92.71	*	+
	HEMBA1006482	71.42	71.44	219.31	199.96	180.73	192.88		
50	HEMBA1006483	2.03	2.96	7.22	3.36	4.88	3.24		
	HEMBA1006485	2.24	1.72	6.31	4.09	6.53	5.51		
	HEMBA1006486	5.08	5.55	13.73	18.4	16	15.46	*	+
55	HEMBA1006489	1.21	2.18	4.1	3.18	5.4	2.17		

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	HEMBA1006158	3.06	4.95	7.5	5.7	9.04	6.31		
	HEMBA1006164	2.61	1.96	5.89	5.26	4.54	5.82		
5	HEMBA1006171	29.76	24.08	54.44	32.3	33.25	34		
	HEMBA1006173	5.15	3.15	30.41	36.16	55.46	57.62	*	+
	HEMBA1006176	315.12	232.27	427.53	476.77	458.01	381.49		
10	HEMBA1006182	1.47	1.71	6.13	3.02	5.84	5.82		
	HEMBA1006197	6.14	5.09	9.26	9.53	12.1	10.41		
	HEMBA1006198	10.07	6.46	26.71	35.03	45.64	55.68	*	+
	HEMBA1006213	1.98	1.78	2.4	2.33	2.69	2.3		
15	HEMBA1006217	44.9	41.62	72.62	94.44	74.92	78.25		
	HEMBA1006226	40.86	36.82	63.27	77.37	79.96	57.39		
	HEMBA1006235	2.13	1.96	4.34	2.6	5.85	2.32		
	HEMBA1006248	1.74	2.03	4.04	3.37	6.4	2.95		
20	HEMBA1006251	5.41	7.29	7.88	11.96	12.95	9.22	*	+
	HEMBA1006252	0.72	1.28	3.67	2.25	3.56	2.92		
	HEMBA1006253	2.13	2.26	4.1	5.93	5.38	5.76	*	+
25	HEMBA1006259	1.96	2.29	6.02	5.7	3.9	5.32		
	HEMBA1006261	12.25	7.27	18.84	14.23	14.08	13.61		
	HEMBA1006268	2.05	2.56	3.86	4.83	5.09	4.06	*	+
	HEMBA1006271	2.04	3.99	10.58	7.13	9.33	8.51		
30	HEMBA1006272	0.97	2.26	2.84	2.38	6.01	1.93		
	HEMBA1006273	1.53	2.09	4.55	3.58	4.52	2.46		
	HEMBA1006276	2.8	1.26	3.62	4.45	5.84	2.82		
	HEMBA1006278	1.57	2.03	3.19	4.08	4.51	2.72		
35	HEMBA1006283	3.09	3	6.08	7.34	11.13	7.09		
	HEMBA1006284	2.47	1.57	3.14	4.75	6.96	2.82		
	HEMBA1006291	1.42	2.56	4.41	4.6	6.16	2.57		
40	HEMBA1006292	3.36	5.12	17.34	19.95	23.83	21.11	*	+
	HEMBA1006293	1.83	1.46	3.19	2.92	4.02	1.36		
	HEMBA1006299	1.92	2.26	7.03	5.02	6.39	4.74		
	HEMBA1006309	2.26	1.43	3.53	4.47	4.69	3.73	*	+
45	HEMBA1006310	4.14	4.32	7.72	9.51	8.34	5.7		
	HEMBA1006311	1.4	2.33	6.68	5.59	5.85	5.72		
	HEMBA1006313	1.2	1.6	2.74	4.29	4.68	2.26		
50	HEMBA1006316	2.16	3.08	6.72	6.63	6.28	5.59		
	HEMBA1006328	2.78	4.28	13.48	15.14	17	16.56		
	HEMBA1006334	1.46	3.1	2.77	2.79	3.95	2.05		
	HEMBA1006335	10.42	13.98	21.66	20.81	18.03	16.77		
55	HEMBA1006344	2.86	3.55	6.9	7.18	6.88	5.75		

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	HEMBA1005999	2.34	4.39	8.84	6.52	8.18	8.75		
	HEMBA1006002	3.53	5	7.73	5.96	10.24	8.99		
5	HEMBA1006005	0.96	2.31	-3.93	2.87	5.65	3.27		
	HEMBA1006011	26.27	24.03	34.08	62.88	44.8	64.92	*	+
	HEMBA1006013	2.43	2.57	6.31	4.17	6.87	3.74		
	HEMBA1006016	1.65	1.98	5.64	3.22	5.34	2.16		
10	HEMBA1006019	2.97	3.23	6.91	4.46	7.22	5.44		
	HEMBA1006021	5.06	6.45	9.21	10.29	12.48	8.77		
	HEMBA1006022	3.19	4.34	6.89	6.52	6.1	5.94		
15	HEMBA1006031	1.32	2.46	4.38	3.72	5.57	5.34		
	HEMBA1006035	3.05	3.72	7.96	4.99	6.37	4.54		
	HEMBA1006036	2.02	2.3	7	4.27	10.14	7.22		
	HEMBA1006042	3.36	3.1	8.51	5.76	9.59	8.31		
20	HEMBA1006044	1.44	1.99	3.61	2.8	3.4	1.67		
	HEMBA1006045	1.98	1.99	5.08	4.3	8.55	5.64		
	HEMBA1006048	2.42	4.18	5.41	6.84	7.96	8.18	*	+
25	HEMBA1006053	1.51	2.72	3.55	3.09	3.3	3.93		
	HEMBA1006055	1.84	1.91	2.46	3.04	4.92	3.67	*	+
	HEMBA1006058	4.04	4.62	11.59	7.42	10.66	12.41		
	HEMBA1006063	9.2	9.36	32.62	26.37	33.39	27.19		
30	HEMBA1006067	4.14	3.27	5.81	5.8	8.4	7.17		
	HEMBA1006081	0.84	2.59	4.77	2.54	7.08	2.49		
	HEMBA1006089	2.58	4.48	6.82	8.28	10.12	9.66	*	+
	HEMBA1006090	1.66	2.31	2.28	1.66	5.26	1.84		
35	HEMBA1006091	1.1	1.35	1.75	3.15	4.31	2.95	*	+
	HEMBA1006093	1.65	1.77	4.21	2.27	6.67	4.26		
	HEMBA1006099	11.9	9.88	23.12	21.57	16.43	20.57		
40	HEMBA1006100	2.78	3.18	13.25	8.38	15.71	13.52		
	HEMBA1006108	2.69	2.08	4.22	3.42	5.24	3.75		
	HEMBA1006114	7.21	8.76	39.36	34.51	62.97	50.98		
	HEMBA1006121	1.18	1.8	3.13	2.43	5.83	2.91		
45	HEMBA1006124	1.79	1.74	4.47	3.11	4.89	4.13		
	HEMBA1006125	18.52	14.19	23.44	25.23	22.56	34.45		
	HEMBA1006130	5.15	3.1	7.57	6.89	7.84	9.51		
	HEMBA1006138	2.43	2.41	5.55	5.27	9.2	8.12		
50	HEMBA1006142	2.62	1.87	6.23	6.67	9.36	8.91	*	+
	HEMBA1006150	8.32	7.44	21.06	20.42	21.82	24.68		
	HEMBA1006151	567.67	524.4	796.07	915.23	875.11	682.15		
55	HEMBA1006155	0.93	1.33	2.92	1.44	5.54	1.94		

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	HEMBA1005749	9.64	9.37	17.99	21.62	19.82	18.15		
	HEMBA1005755	0.49	2.5	3.35	1.66	4.41	2.9		
5	HEMBA1005760	1.59	2.1	-5.37	3.7	4.86	4.14		
	HEMBA1005765	1.47	2.89	5.97	5.91	7.42	7.21		
	HEMBA1005766	8.52	10.49	61.64	93.4	112.09	106.28	*	+
	HEMBA1005780	4.98	5.15	14.49	13.93	16.26	12		
10	HEMBA1005795	1.15	1.8	3.71	4.28	5.45	4.26	*	+
	HEMBA1005809	9.13	9.7	20.95	14.79	29.83	22.75		
	HEMBA1005813	1.76	2.74	6.74	5.39	9	7.08		
15	HEMBA1005815	0.62	1.86	2.75	2.74	4.78	2.13		
	HEMBA1005822	1.34	2.83	5.82	3.77	6.65	5.12		
	HEMBA1005829	1.49	2.66	4.94	5.11	4.53	5.52		
	HEMBA1005833	1.83	1.28	4.41	3.89	5.17	4.57		
20	HEMBA1005834	2.53	1.87	4.84	5.48	11.84	6.81		
	HEMBA1005844	27.89	31.06	52.5	62.42	59.49	58.73	*	+
	HEMBA1005852	5.75	4.18	10.12	10.15	12.34	11.51		
25	HEMBA1005853	2.81	3.05	9.32	5.68	13.84	9.26		
	HEMBA1005878	4.42	4.52	10.14	8.84	10.05	11.01		
	HEMBA1005883	1.67	2.81	4.56	2.81	5.43	4.01		
	HEMBA1005884	1.87	2	3.21	2.27	5.9	2.28		
30	HEMBA1005891	2.39	3.04	3.59	4.34	5.46	3.95		
	HEMBA1005894	2.12	2.41	8.68	8.23	9.08	6.9		
	HEMBA1005898	4.52	3.75	7.4	10.51	10.74	9.53	*	+
35	HEMBA1005902	2.57	3.14	8.63	5.36	6.75	5.82		
	HEMBA1005907	1.15	2.28	2.2	2.28	4.2	1.74		
	HEMBA1005909	0.93	2.68	2.97	2.96	4.55	2.6		
	HEMBA1005911	1.66	3.12	4.9	3.96	9.5	3.41		
40	HEMBA1005912	8.83	7.86	13.57	16.44	8.16	15.58		
	HEMBA1005913	5.05	5.39	10.39	9.57	7.71	7.06		
	HEMBA1005921	2.36	3.86	6.29	5.25	10.06	7.64		
	HEMBA1005922	5.49	5.47	6.99	8.51	10.1	8.74	*	+
45	HEMBA1005929	1.91	2.53	7.53	5.05	5.68	6.47		
	HEMBA1005931	3.32	2.95	6.04	5.31	9.49	6.09		
	HEMBA1005934	2.9	4.69	8.2	8.17	10.14	9.73		
50	HEMBA1005945	3.1	4.12	7.73	6.36	9.39	6.65		
	HEMBA1005962	1.72	1.81	3.16	1.93	3.96	1.72		
	HEMBA1005963	1.86	1.67	3.91	2.26	3.95	2.05		
	HEMBA1005990	6.04	7.39	17.86	14.14	16.18	17.02		
55	HEMBA1005991	2.39	3.35	8.55	7.2	7.64	6.94		

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	HEMBA1005577	1.14	2.02	2.78	0.95	3.08	0.87		
	HEMBA1005581	4.29	4.22	6.85	14.27	18.37	8.15		
5	HEMBA1005582	2.74	2.57	5.35	3.25	7.91	3.21		
	HEMBA1005583	2.47	3.31	5.86	3.32	6.43	3.44		
	HEMBA1005588	2.51	2.85	6.27	5	6.54	6.14		
	HEMBA1005593	1.5	1.4	2.85	1.83	4.57	2.89		
10	HEMBA1005595	2.62	2.82	4.15	3.31	4.65	4.76		
	HEMBA1005597	4.77	5.18	8.13	9.39	10.34	8.64	*	+
	HEMBA1005606	2.29	2.76	5.79	3.96	6.91	5.83		
	HEMBA1005609	2.84	2.64	6.61	4.19	5.77	6.81		
15	HEMBA1005616	2.01	1.66	8.03	5.44	8.75	7.03		
	HEMBA1005621	2.43	1.91	4.42	4.13	6.24	2.7		
	HEMBA1005627	3.84	3.92	11.61	9.73	15.14	14.89		
20	HEMBA1005628	12.1	12.91	20.55	17.92	23.35	18.08		
	HEMBA1005631	13.47	11.94	26.82	22.77	22.87	29.03		
	HEMBA1005632	1.33	2.33	5.06	3.14	3.68	4.47		
	HEMBA1005634	3.06	3.42	5.15	2.81	7.68	5.18		
25	HEMBA1005662	1.18	1.27	3.17	1.06	4.57	3.03		
	HEMBA1005666	5.89	4.51	10.09	10.5	9.01	10.25		
	HEMBA1005670	1	1.08	4.06	2.87	4.35	3.19		
30	HEMBA1005671	2.11	3.38	5.07	5.36	9	5.9		
	HEMBA1005679	2.33	4.64	7.39	6.5	10.44	10.19		
	HEMBA1005680	2.63	2.14	5.9	5.51	7.59	7.72		
	HEMBA1005685	2	1.89	7.27	3.8	6.73	1.97		
35	HEMBA1005698	5.96	4.75	12.88	11.78	14.17	9.93		
	HEMBA1005699	1.4	1	2.45	2.17	3.66	2.96		
	HEMBA1005703	1.22	1.27	3.57	1.79	3.56	1.88		
40	HEMBA1005705	2.39	2.78	6.45	3.41	6.27	3.89		
	HEMBA1005712	1.23	1.34	4.52	2.18	4.84	2.37		
	HEMBA1005717	1.55	1.89	4.7	1.34	5.36	2.16		
	HEMBA1005718	5.27	4.35	7.8	10.09	10	15.72	*	+
45	HEMBA1005721	15.93	20.34	26.12	37.74	25.37	32.79		
	HEMBA1005722	18	19.32	35.72	25.36	30.28	26.91		
	HEMBA1005724	2.17	2.05	4.6	2.47	6.56	2.98		
	HEMBA1005732	1.33	1.54	4.89	7.22	8.38	5.67	*	+
50	HEMBA1005737	1.49	1.19	2.95	2.44	4.57	1.75		
	HEMBA1005742	3.4	4.65	5.7	4.43	6.35	4.22		
	HEMBA1005746	1.2	1.61	3.23	5.69	4.42	4.85	*	+
55	HEMBA1005747	3.8	3.51	7.52	6.08	6.55	6.67		

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	HEMBA1005411	1.38	2.58	4.54	2.84	4.09	2.74		
	HEMBA1005423	4.14	5.87	4.72	12.75	8.59	11.04	*	+
5	HEMBA1005426	0.71	1.79	-2.35	2.08	3.25	4.15		
	HEMBA1005427	9.12	12.47	24.08	37.26	33.57	35.43	*	+
	HEMBA1005430	1.71	1.43	3.23	3.26	4.07	4.78		
	HEMBA1005438	1.42	2.33	4.26	4.01	5.41	2.66		
10	HEMBA1005443	15.57	16.88	35.83	20.61	27.42	29.93		
	HEMBA1005447	1.55	1.95	3.42	2.25	5.9	3		
	HEMBA1005449	1.5	1.36	3.29	1.82	5.83	3.21		
15	HEMBA1005452	8.96	10.79	63.65	107.56	94.42	105.84	*	+
	HEMBA1005454	4.73	4.51	7.91	6.37	8.12	5.33		
	HEMBA1005468	2.49	3.19	4	3.6	4.93	4.86		
	HEMBA1005469	1.39	1.89	5.38	3.44	5.94	4.04		
20	HEMBA1005472	4.04	4.76	7.81	6.79	6.75	8.2		
	HEMBA1005474	4.91	5.42	11.67	7.77	9.4	9.23		
	HEMBA1005475	7.8	7.09	11.75	15.34	10.75	13.59		
25	HEMBA1005489	2.47	3.48	4.94	6.34	7	6.54	*	+
	HEMBA1005497	0.65	1.97	3.47	2.86	4.42	2.14		
	HEMBA1005500	4.64	4.25	11.48	9.17	13.98	10.59		
	HEMBA1005506	3.88	2.15	4.02	2.71	3.57	1.29		
30	HEMBA1005508	5.79	7.67	13.3	11.65	14.52	10.99		
	HEMBA1005511	2.97	2.29	9	4.36	6.36	6.14		
	HEMBA1005513	6.3	9.05	55.46	50.46	60.73	57.45		
35	HEMBA1005517	1.8	2.89	3.88	4.37	6.41	3.74		
	HEMBA1005518	1.48	2.49	3.94	2.94	5.22	2.3		
	HEMBA1005520	1.89	3.19	8.8	8.37	10.13	9.32		
	HEMBA1005522	1.78	2.29	3.37	2.52	4.72	1.45		
40	HEMBA1005526	3.24	2.8	6.25	4.37	7.26	4.58		
	HEMBA1005528	8.59	16.13	16.28	20.24	22.18	20.01		
	HEMBA1005530	2.28	3.62	5.13	4.91	7.75	4.51		
	HEMBA1005538	11.07	10.7	15.34	8.46	4.44	5.89	*	-
45	HEMBA1005539	30.73	31.96	63.32	50.52	49.29	34.26		
	HEMBA1005545	1.2	1.08	3	4.28	6.22	2.25		
	HEMBA1005548	2.38	2.88	11.1	11.04	12.72	9.21		
50	HEMBA1005552	2.76	4.58	12.05	9.13	12.8	10.25		
	HEMBA1005558	1.59	2	5.86	2.18	6.97	4.17		
	HEMBA1005568	3.11	2.96	8.24	6.28	8.45	7.5		
	HEMBA1005570	1.87	2.75	3	1.96	4.49	7.13		
55	HEMBA1005576	1.74	2.8	2.4	3.03	3.55	3.57	*	+

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	HEMBA1005246	8.84	7.84	17.94	12.96	18.21	14.79		
	HEMBA1005251	3.06	2.57	6.81	6.6	8.47	7.24		
5	HEMBA1005252	2.3	2.7	5.31	4.73	11.4	5.01		
	HEMBA1005267	2.77	3.31	7.25	6.77	12.66	9.89		
	HEMBA1005274	1.75	1.53	3.28	4.8	6.38	6.51	**	+
	HEMBA1005275	0.62	1.27	1.59	1.63	3.44	2.12		
10	HEMBA1005288	3.03	1.16	6.1	6.49	6.35	6.68		
	HEMBA1005293	1.46	1.37	2.53	1.5	4.09	1.78		
	HEMBA1005296	440.88	422.67	780.14	598.1	568.79	483.77		
15	HEMBA1005301	3.93	3.26	29.41	28.17	47.57	44.12		
	HEMBA1005304	1.75	2.3	7	3.36	7.29	7.62		
	HEMBA1005305	0.77	1.18	2.57	0.74	3.99	1.71		
	HEMBA1005311	0.97	1.05	2.08	0.81	1.97	2.13		
20	HEMBA1005313	20.35	21.96	39.64	29.02	19.54	26.39		
	HEMBA1005314	1	1.36	2.6	2.2	4.08	1.99		
	HEMBA1005315	0.81	2.41	4.27	1.95	5.32	3.59		
	HEMBA1005317	0.8	1.48	2.93	1.9	3.68	2.43		
25	HEMBA1005318	1.42	0.95	2.68	1.48	4.54	2.21		
	HEMBA1005324	6.24	8.79	28.39	48.36	52.86	57.05	**	+
	HEMBA1005331	2.77	5.64	15.05	21.73	18.68	21.33	*	+
30	HEMBA1005337	12.1	12.25	61.94	98.25	113.65	125.54	*	+
	HEMBA1005338	4.53	4.54	27.15	34.04	35.53	42.4	*	+
	HEMBA1005344	2.75	2.57	5.08	4.17	4.16	3.3		
	HEMBA1005353	1.15	2.78	6.52	4.66	7.21	6.44		
35	HEMBA1005359	2.6	2.82	7.48	6.59	8.03	7.68		
	HEMBA1005362	1.06	2.73	4.93	7.3	11.27	8.37	*	+
	HEMBA1005364	0.96	1.48	3	2.55	3.78	2.75		
	HEMBA1005367	0.96	1.53	4.19	3.3	4.21	2.99		
40	HEMBA1005372	1	2.24	4.18	2.3	5.1	3		
	HEMBA1005374	2.29	3.31	7.83	6.4	6.45	8.09		
	HEMBA1005379	3.19	3.61	7.65	29.57	25.57	24.46	**	+
45	HEMBA1005382	9.85	11.52	79.62	88.72	86.53	120.52		
	HEMBA1005384	1.44	1.76	3.4	2.78	3.74	2.78		
	HEMBA1005386	1.52	2.67	4.54	4.17	4.66	4.84		
	HEMBA1005389	1.1	0.8	3.01	2.51	4.09	3.75		
50	HEMBA1005394	1.64	2.32	5	8.09	10.03	6.99	*	+
	HEMBA1005403	4.19	4.24	8.81	8.46	11.37	8.87		
	HEMBA1005408	1.43	1.71	5.3	4.52	7.12	6.96		
55	HEMBA1005410	1.06	1.74	3.22	1.78	4.97	2.42		

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	HEMBA1005062	2.48	3.06	5.62	2.85	5.13	2.03		
	HEMBA1005066	2.28	2.55	5.72	4.81	8.94	6.89		
5	HEMBA1005067	5.81	10.37	17.44	25.77	30.77	26.84	*	+
	HEMBA1005070	3.33	3.01	8.9	8.51	8.13	6.59		
	HEMBA1005075	1.29	2.45	6.51	3.97	5.38	4.49		
10	HEMBA1005078	7.47	6.74	14.34	14.89	9.95	11.01		
	HEMBA1005079	5.52	5.68	13.65	14.98	16.02	21.42		
	HEMBA1005083	0.94	0.97	2.69	2.17	4.96	1.25		
	HEMBA1005084	5.36	4.49	8.84	7.48	10.38	9.99		
15	HEMBA1005088	1.63	1.64	7.16	3.48	6.18	4.54		
	HEMBA1005089	3.12	3.47	5.53	5.04	9	5.52		
	HEMBA1005090	5.92	5.56	11.7	17.14	12.91	21.06	*	+
	HEMBA1005096	0.88	2.47	3.98	3.35	3.94	2.88		
20	HEMBA1005101	2.29	2.08	4.54	3.23	5.6	4.54		
	HEMBA1005107	1.2	1.92	3.2	2.25	4.48	2.3		
	HEMBA1005113	0.96	2.18	3.35	2.12	5.53	2.72		
25	HEMBA1005123	3.35	3.46	10.83	9.05	10.41	8.28		
	HEMBA1005133	2.6	2.26	7.59	5.11	4.58	7.09		
	HEMBA1005135	1.19	2.77	3.35	1.47	5.28	3.69		
	HEMBA1005145	5.84	6.38	12.56	13.06	14.71	17.84		
30	HEMBA1005149	4.21	3.2	7.92	6.51	8.54	8.47		
	HEMBA1005152	1.81	3.06	3.59	3.31	5.05	3.39		
	HEMBA1005159	1.76	1.96	3.62	2.2	8.12	2.2		
35	HEMBA1005172	120.6	113.48	240.76	174.65	194.57	222.24		
	HEMBA1005185	3.16	2.05	4.47	3.54	6.7	4.15		
	HEMBA1005186	1.55	2.65	3.96	3.52	8.39	2.03		
	HEMBA1005195	2.15	1.08	3.76	1.93	6.73	2.24		
40	HEMBA1005201	3.27	3.45	8.02	5.52	7.17	8.91		
	HEMBA1005202	5.93	4.58	9.4	8.29	9.76	12.83		
	HEMBA1005204	86.99	67.86	141.56	134.68	133.57	172.39		
	HEMBA1005206	6.39	5.08	66.26	72.89	92.07	89.66	*	+
45	HEMBA1005219	4.62	6.56	14.55	15.66	21.85	17.87	*	+
	HEMBA1005223	2.4	1.91	4.92	1.7	3.83	3.86		
	HEMBA1005229	1.18	2.32	3.12	1.57	4.88	2.32		
50	HEMBA1005230	1.22	1.56	4.24	2.71	5.92	4.32		
	HEMBA1005232	0.86	1.13	1.6	0.91	3.81	2.17		
	HEMBA1005238	0.84	1.77	2.67	2.16	3.27	2.29		
	HEMBA1005241	2.77	2.68	7.42	7.63	11.38	7.62		
55	HEMBA1005244	0.87	1.97	5.63	2.57	8.33	6.19		

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	HEMBA1004882	5.8	7.1	10.16	15.5	15.77	12.03	*	+
	HEMBA1004885	2.34	6	8.67	8.58	8.67	9.27		
5	HEMBA1004889	3.25	3.24	6.5	7.67	10.74	9.8	*	+
	HEMBA1004900	1.51	1.59	3.71	3.21	5.14	2.84		
	HEMBA1004909	0.71	1.96	3.59	3.21	4.47	4.97		
	HEMBA1004918	1.46	1.7	4.93	4.03	7.19	5.77		
10	HEMBA1004923	1.1	1.83	3.72	3.98	7.3	4.27		
	HEMBA1004929	1.01	1.7	2.68	1.66	3.59	1.38		
	HEMBA1004930	1.4	2.06	4.31	3.91	5.87	4.72		
15	HEMBA1004933	2.41	1.5	5.32	5.59	5.43	3.38		
	HEMBA1004934	8.61	8.61	17.67	26.2	27.77	26.83	**	+
	HEMBA1004937	1.36	2.42	3.57	2.79	7.51	2.54		
	HEMBA1004943	1.11	1.75	3.63	2.56	4.88	3.68		
20	HEMBA1004944	0.95	2.51	4.73	3.85	4.79	3.58		
	HEMBA1004946	4.78	4.38	12.5	12.69	14.08	12.8		
	HEMBA1004952	0.9	1.83	4.04	2.76	4.41	2		
25	HEMBA1004954	3.55	3.21	7.92	6.02	6.4	6.12		
	HEMBA1004956	1.37	1.11	2.78	2.12	3.66	1.81		
	HEMBA1004960	0.86	0.75	3.27	4.17	6.72	3.06		
	HEMBA1004971	2.41	2.49	6.81	6.45	8.05	6.92		
30	HEMBA1004972	2.57	2.31	3.63	2.07	5.66	5.84		
	HEMBA1004973	1.16	1.78	3.46	2.53	3.27	3.18		
	HEMBA1004977	3.04	3.08	5.36	5.34	6.29	6.36		
	HEMBA1004978	4.53	5.08	14.93	21.43	19.01	25.37	*	+
35	HEMBA1004980	1.92	2.36	7.92	6.55	6.55	7.29		
	HEMBA1004982	0.83	1.36	2.92	2.8	4.45	2.65		
	HEMBA1004983	1.73	1.88	4.29	3.92	5.5	4.21		
40	HEMBA1004995	2.76	3.99	6.53	7.5	7.46	7.41		
	HEMBA1005004	1.63	3.57	5.12	3.45	5.72	4.07		
	HEMBA1005008	1.61	3.52	4.9	4.61	6.42	5.97		
	HEMBA1005009	4.55	3.9	8.37	11.95	6.96	9.22		
45	HEMBA1005019	3.1	2.57	5.78	6.6	8.07	7.19	*	+
	HEMBA1005021	16.12	17.89	30.44	29.81	23.38	23.26		
	HEMBA1005029	3.13	3.42	7.98	7.23	8.66	6.52		
50	HEMBA1005035	6.53	6.29	19.38	18.59	17.77	20.38		
	HEMBA1005036	19.87	20.39	37.72	23.12	19.26	26.98		
	HEMBA1005039	1.7	3.19	4.59	4.36	6.81	3.69		
	HEMBA1005047	4.31	4.06	5.6	7.01	8.54	7.83	**	+
55	HEMBA1005050	2	2.93	5.07	4.33	6.76	4.39		

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	HEMBA1004711	1.22	1.44	5.01	1.19	3.14	3.28		
	HEMBA1004723	4.91	5.18	10.06	11.37	16.92	12.96	*	+
5	HEMBA1004725	4.14	4.9	7.68	5.67	8.62	8.54		
	HEMBA1004730	3.57	2.83	4.74	3.64	8.59	4.41		
	HEMBA1004733	1.07	2.64	2.62	1.06	3.54	2.45		
	HEMBA1004734	2.57	3.22	4.46	2.68	5.28	5.32		
10	HEMBA1004736	1.1	1.89	7.12	3.87	12.1	5.09		
	HEMBA1004748	2.24	0.94	5.6	3.02	5.22	2.83		
	HEMBA1004749	6.88	8.33	19.68	16.49	19.03	23.29		
15	HEMBA1004751	1.96	1.76	5.55	3.99	9.86	5.04		
	HEMBA1004752	1.51	1.6	4.23	4.56	4.11	3.32		
	HEMBA1004753	29.15	25.19	85.53	59.89	62.5	95.58		
	HEMBA1004755	7.02	6.32	12.37	9.73	12.72	14.63		
20	HEMBA1004756	1.45	1.76	3.86	2.34	5.21	2.17		
	HEMBA1004758	1.18	1.64	4.53	3.92	5.65	3.17		
	HEMBA1004763	1.79	2.39	5.56	5.45	6.53	6.09		
25	HEMBA1004768	0.83	1.64	2.89	1.69	4.26	1.38		
	HEMBA1004770	1.09	1.36	2.43	1.47	3.53	1.94		
	HEMBA1004771	0.99	1.02	2.44	2.18	3.26	2.57		
	HEMBA1004775	4.07	3.84	7.29	8.61	9.62	13.74	*	+
30	HEMBA1004776	1.86	3.21	3.33	3.95	6.83	6.04	*	+
	HEMBA1004778	1.75	2.24	6.11	3.21	7.64	5.74		
	HEMBA1004784	1.51	1.59	3.11	2.36	4.18	3.62		
	HEMBA1004785	1	1.78	4.15	2.18	6.54	4.48		
35	HEMBA1004789	2.34	2.07	4.42	1.87	5.64	2.87		
	HEMBA1004795	0.62	1.89	3.13	2.45	4.23	1.42		
	HEMBA1004797	1.06	0.84	1.85	2.31	2.85	2.76	*	+
40	HEMBA1004803	4.98	1.72	5.31	4.62	9.79	4.82		
	HEMBA1004806	1.23	1.78	3.22	2.36	3.83	2.33		
	HEMBA1004807	3.05	1.95	3.86	5.58	7.44	4.14		
	HEMBA1004816	4.73	2.61	3.59	3.78	7.97	9.34		
45	HEMBA1004820	1.73	2.33	4.6	2.97	8.74	3.71		
	HEMBA1004833	1.22	1.23	5.54	2.71	6.95	4.37		
	HEMBA1004847	4.73	2.8	9.84	8	11.83	9.15		
	HEMBA1004850	1.01	1.78	6.15	6.56	13.54	6.38		
50	HEMBA1004863	1.75	2.3	4.92	3.34	4.45	5.17		
	HEMBA1004864	2.66	3.91	6.68	4.19	9.03	6.51		
	HEMBA1004865	1.13	2.61	4.06	3.86	6.64	3.38		
55	HEMBA1004880	2.22	3.32	9.22	9.9	10.38	9.48		

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	HEMBA1004554	1.21	1.05	2.53	1.91	7.29	2.16		
	HEMBA1004558	2.98	4.6	12.99	21.19	21.7	23.66	**	+
5	HEMBA1004560	1.74	2.06	4.16	3.47	6.18	4.25		
	HEMBA1004564	4.31	3.3	8.73	7.27	9.96	8.65		
	HEMBA1004566	50.51	49.26	83.43	126.94	55.59	138.13		
	HEMBA1004573	1.07	1.97	3.91	2.91	4.11	2.3		
10	HEMBA1004576	3.68	3.22	10.49	4.73	5.64	3.11		
	HEMBA1004577	6.11	7.2	14	11.74	13.04	10.71		
	HEMBA1004586	1.56	1.32	5.09	5.01	6.65	4.67		
15	HEMBA1004596	3.97	3.19	19.1	27.4	39.04	32.16	*	+
	HEMBA1004604	9.04	8.14	40.08	64.86	69.99	62.02	*	+
	HEMBA1004607	0.96	1.97	4.17	3.02	4.39	3.96		
	HEMBA1004610	1.16	1.89	3.19	2.91	4.8	2.64		
20	HEMBA1004617	0.78	0.89	3.29	2.77	4.53	2.51		
	HEMBA1004622	1.05	2.39	5.1	4.07	5.84	5.04		
	HEMBA1004626	1.56	2.36	4.75	4.06	4.93	2.33		
25	HEMBA1004629	1.5	1.34	3.95	3.17	4.58	2.74		
	HEMBA1004631	1.57	1.73	4.41	3.3	5.73	4.97		
	HEMBA1004632	1.02	1.3	3.43	2.38	3.64	2.36		
	HEMBA1004633	3.2	3.98	8.84	9.3	9.04	10.74		
30	HEMBA1004636	1.29	2.07	3.22	3.23	3.61	6.58		
	HEMBA1004637	1.57	2.12	4.19	3.97	4.85	2.39		
	HEMBA1004638	1.31	1.67	3.26	1.8	4.24	1.52		
	HEMBA1004645	3.04	2.88	6.5	5.07	6.21	5.08		
35	HEMBA1004656	4.38	2.76	4.96	4.13	4.58	3.39		
	HEMBA1004657	16.78	17.12	35.48	31.85	17.55	22.74		
	HEMBA1004666	1.27	2.2	3.32	2.78	4.52	1.44		
40	HEMBA1004669	2.49	3.6	6.16	7.32	8.34	5.55		
	HEMBA1004670	3.1	2.74	6.27	5.11	6.96	3.3		
	HEMBA1004672	1.29	2.33	4.85	2.58	7.25	3.63		
	HEMBA1004689	23.54	21.34	82.29	90.21	98.37	106.84		
45	HEMBA1004690	4.74	5.24	15.26	19.89	22.39	20.76	*	+
	HEMBA1004693	3.16	5.98	25.39	24.92	35.74	32.68		
	HEMBA1004697	1.64	1.96	4.88	7.14	9.69	4.03		
50	HEMBA1004702	8.73	11.47	19.57	19.47	14.63	11.2		
	HEMBA1004704	1.9	3.35	9.01	3.41	5.17	5.36		
	HEMBA1004705	1.13	1.93	3.32	1.8	4.34	1.54		
	HEMBA1004706	1.34	2.4	3.89	4.31	6.23	4.7		
55	HEMBA1004709	2.96	2.9	10.14	6.81	8.65	6.93		

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	HEMBA1004366	1. 97	1. 91	3. 57	4. 53	6. 9	4. 73	*	+
	HEMBA1004372	0. 3	0. 67	1. 03	2. 04	1. 85	0. 96		
5	HEMBA1004377	6. 57	5. 58	-10. 48	16. 45	11. 58	13. 72	*	+
	HEMBA1004389	8. 39	7. 87	13. 69	16. 87	19. 3	11. 15		
	HEMBA1004391	1. 18	0. 88	1. 37	1. 27	4. 23	2. 34		
	HEMBA1004393	65. 85	84. 9	114. 17	77. 61	53. 19	85. 8		
10	HEMBA1004394	0. 84	1. 14	1. 88	1. 68	3. 9	2. 59		
	HEMBA1004396	1. 62	0. 82	3. 24	2. 43	5. 44	2. 15		
	HEMBA1004401	4. 33	4. 37	5. 25	8. 95	14. 42	11. 05	*	+
15	HEMBA1004405	3. 86	2. 57	5. 7	7. 45	7. 46	9. 38	*	+
	HEMBA1004408	4. 27	2. 66	5. 34	7. 19	8. 8	9. 85	*	+
	HEMBA1004414	1. 72	1. 74	5. 93	9. 48	8. 98	13. 19	*	+
	HEMBA1004429	2. 95	3. 96	4. 23	4. 68	6. 9	6. 8	*	+
20	HEMBA1004433	1. 27	1. 43	2. 98	2. 55	2. 65	3. 47		
	HEMBA1004440	1. 33	1. 33	2. 62	2. 3	2. 63	1. 79		
	HEMBA1004444	2. 73	1. 9	4. 47	4. 64	5. 38	4. 66		
	HEMBA1004446	1. 37	0. 95	2. 26	1. 84	2. 94	3. 19		
25	HEMBA1004451	3. 79	7. 37	7. 66	12. 15	9. 81	9. 87	*	+
	HEMBA1004452	0. 71	1. 96	2. 89	3. 53	8. 91	2. 36		
	HEMBA1004454	1. 56	1. 71	3. 06	2. 81	4. 83	4. 1		
30	HEMBA1004460	1. 91	3. 49	7. 49	4. 39	6. 91	5. 19		
	HEMBA1004461	1	1. 87	1. 43	1. 09	3. 95	1. 99		
	HEMBA1004468	3. 22	4. 71	7. 36	9. 92	12. 73	8. 79	*	+
	HEMBA1004479	1. 02	1. 38	2. 44	2. 59	5. 99	3. 42		
35	HEMBA1004482	2. 77	1. 93	5. 5	5. 62	5. 83	5. 6		
	HEMBA1004491	6. 18	5. 55	13	12. 32	15. 5	15. 72		
	HEMBA1004499	9. 09	12. 56	65. 85	86. 22	91. 62	102. 95	*	+
40	HEMBA1004502	1. 81	1. 9	4. 82	3. 14	4. 83	2. 49		
	HEMBA1004505	1. 57	2. 42	4. 6	4. 78	7. 66	4. 09		
	HEMBA1004506	0. 96	2. 32	3. 48	3. 28	5. 58	2. 73		
	HEMBA1004507	29. 33	30. 94	74. 36	85. 26	112. 08	83. 25	*	+
45	HEMBA1004509	1. 62	2. 92	3. 8	5. 09	5. 19	4. 5	*	+
	HEMBA1004523	1. 04	1. 68	1. 65	1. 82	3. 64	3. 02		
	HEMBA1004528	9. 41	10. 46	59. 94	86. 06	89. 35	98. 27	*	+
50	HEMBA1004534	6. 43	9. 06	20. 99	14. 78	16. 24	20. 46		
	HEMBA1004536	1. 91	1. 91	4. 04	2. 58	5. 65	4. 53		
	HEMBA1004538	10. 84	12. 11	16. 22	14. 06	12. 69	14. 98		
	HEMBA1004542	4. 37	3. 88	23. 08	31. 45	41. 94	35. 14	*	+
55	HEMBA1004552	2. 98	1. 35	4. 27	9. 89	9. 44	7. 1	**	+

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	HEMBA1004243	1.8	2.09	3.58	2.8	3.03	2.76		
	HEMBA1004246	1.6	2.68	5.65	6.18	6.24	6.15		
5	HEMBA1004247	0.89	2.73	3.74	3.69	4.23	3.37		
	HEMBA1004248	4.01	3.54	3.85	5.91	8.31	7.47	**	+
	HEMBA1004250	1.55	2.16	2.87	1.91	5.22	1.47		
	HEMBA1004252	3.57	3.27	4.8	4.64	5.79	4.28		
10	HEMBA1004260	2.56	3.08	6.87	7.32	8.16	7.61		
	HEMBA1004264	1.26	2.11	2.59	2.16	2.86	1.37		
	HEMBA1004267	5.5	5.81	14.29	14.22	12.19	11.57		
15	HEMBA1004272	1.75	2.31	3.31	2.26	3.84	2.04		
	HEMBA1004274	5.83	8.13	58.69	77.19	87.61	76.22	*	+
	HEMBA1004275	1	5.4	3.34	1.49	4.49	2.42		
	HEMBA1004276	2.27	2.2	3.42	3.45	4.2	3.06		
20	HEMBA1004279	2.13	2.33	4.37	3.29	5.2	3.88		
	HEMBA1004284	1.78	2.56	6.03	4.16	4.9	5.23		
	HEMBA1004286	1.41	1.35	2.44	1.65	3.55	2.1		
25	HEMBA1004289	2.58	4.17	5.59	5.16	7.18	8.44		
	HEMBA1004293	20.24	18.64	51.03	77.3	52.39	74.25	*	+
	HEMBA1004295	1.08	2.65	3.08	2.73	4.02	2.05		
	HEMBA1004302	0.72	1.84	2.29	1.21	3.39	1.49		
30	HEMBA1004306	2.11	3.01	5.96	3.99	5.74	6.01		
	HEMBA1004312	1.3	1.58	4.98	3.57	3.56	3.56		
	HEMBA1004314	1.78	1.86	4.1	3.35	6.23	4.38		
	HEMBA1004321	0.88	1.66	2.56	4.05	4.15	4.82	**	+
35	HEMBA1004323	2.4	3.16	4.7	4.77	5.29	5.1		
	HEMBA1004327	1.18	1.78	3.23	3.51	4.46	3.1		
	HEMBA1004329	5.57	6.73	15.22	16.29	15.33	15.14		
40	HEMBA1004330	3.93	3.54	4.06	5.87	7.54	5.36	*	+
	HEMBA1004334	2.92	3.63	4.69	4.18	4.74	6.06		
	HEMBA1004335	1.15	1.77	5.1	3.04	4.75	3.14		
	HEMBA1004341	1.01	1.1	1.25	1.13	3.34	1.14		
45	HEMBA1004344	29.93	34.05	68.18	74.25	94.77	82.54	*	+
	HEMBA1004347	0.67	1.65	2.21	2.31	2.25	2.78		
	HEMBA1004349	12.07	11.55	22.26	19.39	22.71	24.41		
50	HEMBA1004352	2.06	2.56	6.96	6.05	6.92	6.58		
	HEMBA1004353	10.21	14.95	25.3	19.64	26.63	25.83		
	HEMBA1004354	1.9	2.56	6.29	5.42	6.94	5.83		
	HEMBA1004356	5.75	6.89	9.43	20.06	19.7	20.1	**	+
55	HEMBA1004360	1.35	1.16	2.73	1.23	4.23	2.26		

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	HEMBA1004067	10.12	14.76	90.67	108.89	125.21	128.6	*	+
	HEMBA1004071	7.51	7.77	16.52	17.31	12.23	13.37		
5	HEMBA1004074	0.78	1.93	-3.97	4.48	7.06	5.69	*	+
	HEMBA1004078	3.87	2.95	5.22	6.52	6.2	6.87	*	+
	HEMBA1004085	1.05	1.19	2.83	3.57	4.57	2.45		
	HEMBA1004086	3.38	4.95	6	8.92	8.09	6.51	*	+
10	HEMBA1004097	1.18	1.13	2.97	3.66	3.28	2.97		
	HEMBA1004100	3.85	4.81	8.96	6.9	9.64	9.55		
	HEMBA1004103	2	2.91	6.25	6.25	7.24	7.38		
15	HEMBA1004110	3	3.77	5.43	4.18	4.23	5.02		
	HEMBA1004111	3.96	7.64	44.2	53.81	60.1	57.3	*	+
	HEMBA1004124	7.14	10.51	60.12	83.27	97.96	83.59	*	+
	HEMBA1004130	3.12	3.46	10.29	9.45	6.84	8.43		
20	HEMBA1004131	2.14	2.12	3.06	4.08	3.73	3.21	*	+
	HEMBA1004132	0.77	2.22	4.84	3.94	6.31	4.2		
	HEMBA1004133	0.69	1.77	2.56	3.28	5.17	3.22		
25	HEMBA1004138	0.89	1.19	3.05	2.21	4.11	1.83		
	HEMBA1004143	7.1	7.48	17.43	18.83	15	17.6		
	HEMBA1004146	0.89	2.03	3.01	2.96	4.21	2.69		
	HEMBA1004148	1.85	1.57	2.13	2.25	3.38	1.99		
30	HEMBA1004149	1.54	1.44	2.77	2.83	2.59	3.32		
	HEMBA1004150	0.49	1.06	2.15	2.31	1.58	1.08		
	HEMBA1004154	2.24	1.64	5.28	6.28	7.07	4.61		
35	HEMBA1004164	1.84	2.23	5.63	6.89	7.13	5.81		
	HEMBA1004168	2.16	2.24	4.69	3.9	5.32	7.84		
	HEMBA1004199	1.37	1.92	2.34	3.17	3.66	1.8		
	HEMBA1004200	0.84	1.98	3	1.5	4.05	1.78		
40	HEMBA1004201	4.87	5.68	17.64	26.94	32.17	25.65	*	+
	HEMBA1004202	7.7	10.5	9.9	18.08	16.29	15.77	**	+
	HEMBA1004203	1.63	2.31	3.66	4.5	5.3	4.44	*	+
	HEMBA1004207	1.9	3.24	3.62	5.73	6.23	6.2	**	+
45	HEMBA1004210	1.13	1.72	2.67	1.95	4.14	1.87		
	HEMBA1004225	1.1	2.47	5.23	5.96	7.12	5.4		
	HEMBA1004227	2.17	4.44	3.86	5.14	5.71	5.16		
50	HEMBA1004235	2.68	2.91	3.74	5.79	5.78	4.44	*	+
	HEMBA1004237	3	3.31	5.23	5.95	4.67	5.47		
	HEMBA1004238	2.06	3.24	5.93	5.84	7.64	6.52		
	HEMBA1004241	2.32	3.09	3.87	2.74	3.74	3.35		
55	HEMBA1004242	8.66	13.05	20.15	26.83	32.28	26.48	*	+

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	HEMBA1003945	12.57	13.75	22.75	20.99	14.77	19.74		
	HEMBA1003949	1.4	1.9	3.53	3.29	6.22	4.14		
5	HEMBA1003950	3.46	4.86	-6.49	14.69	17.53	13.02	**	+
	HEMBA1003953	1.91	1.6	5.14	0.72	3.97	1.44		
	HEMBA1003958	5.16	3.6	7.47	7.54	9.45	6.64		
	HEMBA1003959	2.42	2.72	5.72	5.5	5.5	9.02		
10	HEMBA1003960	3.25	5.81	34.7	24.04	26.4	28.28		
	HEMBA1003966	9.63	8.28	16.73	16.75	17.67	19.84		
	HEMBA1003967	1.75	3.06	3.47	3.48	3.6	3.27		
	HEMBA1003968	0.97	2.14	2.55	2.49	4.56	1.82		
15	HEMBA1003974	634.2	699.64	821.36	986.23	1340.97	1248.21	*	+
	HEMBA1003976	1.05	1.84	3.36	1.21	3.27	2.04		
	HEMBA1003977	1.48	2.07	1.99	1.41	3.49	2.15		
20	HEMBA1003978	2.91	3.72	3.54	3.77	6.18	3.53		
	HEMBA1003981	9.01	6.77	14.06	12.05	11.49	18.27		
	HEMBA1003982	102.64	103.61	302.15	380.08	375.9	466.69	*	+
25	HEMBA1003985	1.18	1.9	2.43	3.21	3.79	2.18		
	HEMBA1003987	3.04	2.23	3.1	2.56	4.34	5.53		
	HEMBA1003989	1.62	1.77	4.56	3.79	5.12	3.31		
	HEMBA1004000	1.63	2.35	5.05	3.46	5.35	4.18		
30	HEMBA1004006	2.79	2.88	12.86	16.29	22.13	19.73	*	+
	HEMBA1004007	0.7	1.92	5.28	3.03	5.18	4.72		
	HEMBA1004010	67.4	61.25	98.24	112.56	96.78	136.86		
	HEMBA1004011	0.48	1.74	2.18	2.58	3.29	1.62		
35	HEMBA1004012	0.79	1.84	2.3	3.11	4.8	3.53	*	+
	HEMBA1004015	2.68	4.15	5.38	8.68	10.65	9.21	**	+
	HEMBA1004024	1.47	2.73	5.65	5.68	8.26	8.07		
40	HEMBA1004029	1.93	3.1	3.03	4.6	8.38	13.11		
	HEMBA1004038	1.04	1.24	1.55	1.18	3.38	1.59		
	HEMBA1004042	0.89	1.42	2.22	1.58	4	2.48		
	HEMBA1004045	0.28	0.94	2.42	3.07	3.32	2.53		
45	HEMBA1004048	4.16	4.16	12.1	19.93	14.84	22.3	*	+
	HEMBA1004049	3.56	3.18	4.87	4.92	6.83	5.48		
	HEMBA1004051	136.19	118.77	205.49	243.62	283.22	223.29	*	+
50	HEMBA1004053	5.11	4.64	8.92	25.25	27.24	21	**	+
	HEMBA1004055	2.28	3.2	4.24	2.15	5.51	2.86		
	HEMBA1004056	3.78	3.07	6.73	5.3	10.99	9.56		
	HEMBA1004060	0.86	1	1.7	0.78	3.94	1.65		
55	HEMBA1004061	4.76	3.94	6.44	7.37	12.64	8.57		

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	HEMBA1003764	0.88	2.06	4.9	1.86	4.36	4.24		
	HEMBA1003769	6.61	8.95	15.57	22.58	17.05	20.77	*	+
5	HEMBA1003773	2.16	3.5	4.48	5.8	6.7	5.98	*	+
	HEMBA1003783	3.12	3.11	4.95	8.58	8.64	8.27	**	+
	HEMBA1003784	0.46	1.37	2.89	2.37	2.45	1.77		
	HEMBA1003794	3.48	3.61	13.64	11.98	11.61	15.55		
10	HEMBA1003799	1.09	1.05	4.29	2.44	5.06	3.76		
	HEMBA1003803	7.58	6.67	12.05	11.68	7.64	8.41		
	HEMBA1003804	1.08	2.53	4.45	1.86	3.35	2.05		
15	HEMBA1003805	6.41	7.87	10.45	15.74	7.09	11.93		
	HEMBA1003807	1.52	1.53	3.21	2.71	6.32	2.37		
	HEMBA1003810	1.72	3.29	6.06	5.98	4.85	4.55		
	HEMBA1003827	2.71	4.55	12.08	13.28	10.48	15.35		
20	HEMBA1003836	3.42	4.84	10.27	11.16	12.81	9.96		
	HEMBA1003838	16.58	16.15	31.32	34.24	33.25	35.46		
	HEMBA1003843	4.6	6.54	7.01	13.61	6.48	11.42		
25	HEMBA1003846	19.54	21.94	61.32	72.86	70.58	83.4	*	+
	HEMBA1003856	1.41	1.66	2.85	2.07	4.03	2.51		
	HEMBA1003857	2.89	3.1	5.85	5.89	8.29	6.88		
	HEMBA1003864	1.56	2.61	4.04	3.32	4.03	2.75		
30	HEMBA1003866	0.89	0.75	2.21	1.66	2.23	0.73		
	HEMBA1003868	10.92	10.88	18.59	13.26	7.59	15.72		
	HEMBA1003879	0.95	1.33	3.16	3.49	4.42	3.09		
	HEMBA1003880	1.81	2.35	2.78	3.53	4.78	2.3		
35	HEMBA1003884	10.97	11.37	39.03	54.69	62.46	57.8	*	+
	HEMBA1003885	4.59	4.82	7.14	9.19	6.32	8.41		
	HEMBA1003887	3.58	4.93	7.7	8.65	7.93	8.18		
40	HEMBA1003890	4.2	4.48	7.18	7.53	9.1	6.26		
	HEMBA1003893	4.38	6.39	9.53	8.75	13.24	9.94		
	HEMBA1003896	4.15	4.15	10.62	7.4	9.12	6.43		
	HEMBA1003902	1.39	3.78	5.09	4.91	6.42	5.1		
45	HEMBA1003904	0.87	2.16	2.46	2.82	4.32	2.11		
	HEMBA1003908	1.18	1.3	2.89	2.12	5.25	1.43		
	HEMBA1003926	14.46	12.2	39.79	45.5	34.97	55.56		
50	HEMBA1003937	2.75	3.31	5.38	4.3	6.85	4.57		
	HEMBA1003939	2.43	2.48	6.56	8.3	13.32	8.04		
	HEMBA1003940	2.45	3.08	5.01	4.29	6.22	5.55		
	HEMBA1003941	1.4	2.26	2.48	3.37	4.57	4.42	*	+
55	HEMBA1003942	1.63	2.88	3.13	2.01	3.85	2.22		

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	HEMBA1003610	2.33	3.2	4.48	6.12	5.64	6.81	*	+
	HEMBA1003615	1.76	2.61	5.23	4.95	5.21	4.96		
5	HEMBA1003617	3.59	3.54	8.59	6.92	11.37	8.5		
	HEMBA1003620	5.76	6.01	4.98	13.48	17.69	12.58	**	+
	HEMBA1003621	1.6	1.66	3.19	4.52	5.42	5.08	**	+
	HEMBA1003622	0.96	0.69	1.38	1.47	3.17	2.25		
10	HEMBA1003630	0.78	1.02	1.95	1.68	2.97	1.55		
	HEMBA1003637	0.66	1.93	2.59	2.11	3.11	2.63		
	HEMBA1003640	2.33	2.1	5.27	4.16	5.68	5.5		
15	HEMBA1003645	1.12	1.2	4.41	2.3	3.82	3.06		
	HEMBA1003646	0.94	1.21	1.76	1.25	3.25	1.8		
	HEMBA1003647	0.49	2.15	3.27	2.46	3.79	2.21		
	HEMBA1003656	3.32	3.77	6.96	17.01	10.45	13.78	*	+
20	HEMBA1003662	1.37	2.08	1.54	5.2	3.81	4.91	**	+
	HEMBA1003666	23.84	17.7	51.57	21.97	21.85	24.71		
	HEMBA1003667	4.74	3.63	6.03	4.61	6.22	7.09		
	HEMBA1003670	0.83	0.65	1.94	1.18	2.61	1.51		
25	HEMBA1003674	32.16	29.41	63.99	118.95	138.25	123.17	**	+
	HEMBA1003677	1.84	2.06	4.28	2.32	5.31	3.78		
	HEMBA1003679	1.2	1.68	3.72	2.22	6.19	3.23		
30	HEMBA1003680	4.55	4.68	20.52	27.26	28.13	28.07	*	+
	HEMBA1003684	1.57	1.9	3.98	4	3.65	4.47		
	HEMBA1003690	6.22	7.41	8.65	7.94	9.93	7.33		
	HEMBA1003692	2.41	3.82	7.23	8	8.28	7.7		
35	HEMBA1003702	2.64	3.82	4.83	7.11	6.86	6.07	*	+
	HEMBA1003711	1.06	1.21	3.39	2.93	3.88	2.37		
	HEMBA1003714	1.31	1.26	2.13	1.61	2.45	1.42		
40	HEMBA1003715	1.46	2.7	6.58	10.21	9.15	6.87	*	+
	HEMBA1003717	1.91	2.31	3.91	3.03	3.66	4.38		
	HEMBA1003720	0.81	2.6	5.07	4.16	4.16	4.21		
	HEMBA1003725	0.83	1.57	2.47	3.22	4.91	3.17	*	+
45	HEMBA1003728	1.28	2.48	3.4	2.65	4.36	2.72		
	HEMBA1003729	0.98	2.35	2.85	3.6	4.36	3.52	*	+
	HEMBA1003732	1.11	1.52	3.49	3.01	2.75	1.88		
	HEMBA1003733	1.18	1.9	2.94	3.7	4.95	3.92	*	+
50	HEMBA1003742	5.15	7.3	5.95	21.53	22.58	19.56	**	+
	HEMBA1003743	1.37	1.76	3.21	4.13	4.36	3.68	*	+
	HEMBA1003758	3.26	3.29	11.72	10.07	16.03	12.24		
55	HEMBA1003760	0.82	2.43	3.09	1.92	4.19	3.16		

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	HEMBA1003447	7.55	9.08	49.72	65.41	63.46	65.15	*	+
	HEMBA1003453	21.03	22.03	42.15	27.85	29.02	27.64		
5	HEMBA1003461	1.5	2.13	3.49	2.6	3.63	2.2		
	HEMBA1003463	2.82	3.68	6.02	5.97	3.84	6.41		
	HEMBA1003465	1.77	2.21	6.31	4.75	5.02	3.82		
	HEMBA1003480	2.58	3.91	8.62	9.63	9.6	9.42		
10	HEMBA1003485	7.06	4.84	5.29	6.13	7.26	5.52		
	HEMBA1003487	1.8	1.85	3.4	7.12	6.39	6.79	**	+
	HEMBA1003492	1.42	1.95	4.11	2.41	5.87	2.1		
15	HEMBA1003494	9.36	8.61	12.16	18.24	18.69	17.83	**	+
	HEMBA1003497	2.19	2.16	3.29	3.35	6.06	2.97		
	HEMBA1003503	0.98	1.74	3.37	5.04	3.18	2.13		
	HEMBA1003511	0.99	2.19	3.7	2.3	4.42	2.5		
20	HEMBA1003528	3.33	4	6.51	5.77	5.04	4.46		
	HEMBA1003530	1.33	0.85	3.62	1.97	3.15	2.45		
	HEMBA1003531	1.14	1.72	5.39	4.74	7.24	4.51		
25	HEMBA1003532	12.97	14.66	34.3	28.69	25.31	31.26		
	HEMBA1003538	2.54	2.4	17.88	14.54	21.58	16.83		
	HEMBA1003545	0.68	2.08	3.17	1.85	3.6	2.17		
	HEMBA1003546	1.27	2.03	1.68	1.98	2.15	2.42		
30	HEMBA1003548	1.4	3.18	3.6	1.41	4.15	2.23		
	HEMBA1003553	31.29	31.45	47.99	54.36	41.34	45.65		
	HEMBA1003555	1.39	2.73	4.81	3.53	4.48	5.19		
35	HEMBA1003556	1.24	1.76	2.96	3.14	5.75	3.31		
	HEMBA1003560	1.89	2.66	7.87	10.08	13.24	9.9	*	+
	HEMBA1003565	54.27	66.88	96.28	121.29	139.88	148.68	*	+
	HEMBA1003568	1.86	2.27	3.24	2.36	7.41	2.78		
40	HEMBA1003569	2.93	2.61	2.96	5.07	3.95	4.53	**	+
	HEMBA1003571	3.53	2.33	3.8	5.19	5.3	5.83	*	+
	HEMBA1003579	3.51	4.29	4.83	3.79	5.68	5.91		
	HEMBA1003580	3.82	4.09	4.96	3.11	4.41	3.53		
45	HEMBA1003581	0.82	2.62	2.07	1.63	3.19	2.4		
	HEMBA1003591	10.8	11.44	30.24	33.74	35.7	36.88	*	+
	HEMBA1003595	0.93	1.16	2.46	2.98	4.02	2.01		
50	HEMBA1003597	3.15	3.18	8.74	10.82	11.39	11.59	*	+
	HEMBA1003598	0.58	0.93	1.33	2.62	1.83	1.61	*	+
	HEMBA1003600	3.71	4.19	13.35	14.77	13.86	16.69		
	HEMBA1003602	2.84	2.64	4.89	5.89	6.97	9.14	*	+
55	HEMBA1003604	2.3	3.35	5.67	6.63	8.29	8.16	*	+

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	HEMBA1003281	1.83	1.29	3.53	1.79	3.85	2.48		
	HEMBA1003284	1.24	1.91	3.43	3.03	5.05	3.32		
5	HEMBA1003286	7.75	6.73	34.23	45.25	71.61	51.97	*	+
	HEMBA1003291	1.65	1.91	4.84	3.21	3.32	3.25		
	HEMBA1003294	1.89	3.5	7.47	4.86	5.7	5.62		
	HEMBA1003296	4.74	8.32	46.61	63.82	70.23	54.45	*	+
10	HEMBA1003304	0.77	1.44	2.88	2.91	5.37	1.87		
	HEMBA1003306	4.37	6.3	10.28	15.7	17.76	11.56	*	+
	HEMBA1003309	0.91	1.9	2.85	2.87	4	2.3		
	HEMBA1003314	1.43	2.26	3.82	4.48	3.52	4.02		
15	HEMBA1003315	6.37	4.38	10.14	15.2	16.23	17.88	**	+
	HEMBA1003322	4.81	5.92	10.9	8.46	10.83	8.07		
	HEMBA1003326	1.94	3.97	5.55	2.93	7.4	3.68		
20	HEMBA1003327	0.81	1.61	3.63	2.36	4.3	2.28		
	HEMBA1003328	0.76	2.43	5.38	4.25	5.51	5.06		
	HEMBA1003330	2.27	2.81	4.84	4.66	5.83	6.94		
	HEMBA1003348	3.22	2.45	11.3	11.28	13.98	16.37		
25	HEMBA1003369	2.39	2.6	7	9.64	8.65	5.33		
	HEMBA1003370	3.14	3.6	8.85	12.54	10.83	13.98	*	+
	HEMBA1003373	1.12	1.3	3.4	2.14	5.05	2.94		
30	HEMBA1003376	3.75	2.83	7.71	9.83	12.46	10.39	*	+
	HEMBA1003380	1.12	2.3	3.63	2.25	3.9	2.57		
	HEMBA1003384	0.98	1.71	2.91	2.11	4.78	2.14		
	HEMBA1003387	1.3	1.24	2.14	1.83	3.24	1.98		
35	HEMBA1003392	2.51	2.28	3.43	5.21	5.91	4.44	*	+
	HEMBA1003395	1.02	1.45	2.84	4.06	4.29	2.18		
	HEMBA1003399	1.03	1.4	3.27	3.21	3.26	2.19		
	HEMBA1003400	1.36	2.22	4.64	3.23	7.19	5.22		
40	HEMBA1003402	1.62	1.74	3.29	2.32	4.22	2.59		
	HEMBA1003403	7.13	9.32	50.9	66.1	66.81	77.49	*	+
	HEMBA1003408	3.68	4.5	7.27	6.02	5.77	7.71		
45	HEMBA1003412	5.08	6.79	8.35	10.96	8.79	9.75		
	HEMBA1003417	5.71	6.5	10.15	8.18	8.86	7.36		
	HEMBA1003418	4.01	5.12	6.53	7.37	11.45	9.3	*	+
	HEMBA1003420	16.29	17.91	35.46	33.32	34.37	32.89		
50	HEMBA1003425	0.76	1.65	3.06	2.33	3.58	2.21		
	HEMBA1003433	1.4	2.43	3.34	4.88	4.54	4.09	*	+
	HEMBA1003440	11.39	12.08	19.86	24.13	13.99	24.26		
55	HEMBA1003442	4.37	4.67	4.94	3.54	6.73	5.96		

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	HEMBA1003117	2.1	3.41	4.4	3.36	4.99	2.44		
	HEMBA1003120	3.02	2.65	5.55	3.23	7.38	4.29		
5	HEMBA1003129	2.47	2.6	-6.66	10.28	6.19	7.28		
	HEMBA1003133	2.05	4.74	7.61	7.74	7.59	5.59		
	HEMBA1003136	2.64	3.59	5.25	5.37	5.88	4.49		
	HEMBA1003142	2.01	2.27	6.15	6.62	6.35	5.34		
10	HEMBA1003148	1.3	1.4	2.82	1.49	4.26	1.6		
	HEMBA1003151	1.91	2.08	4.23	2.9	5.34	4.24		
	HEMBA1003152	3.27	1.98	5.84	4.74	5.71	2.58		
15	HEMBA1003157	1.23	1.88	2.58	4.2	5.38	3.21	*	+
	HEMBA1003166	6.14	6.06	14.06	22.98	18.03	21.74	*	+
	HEMBA1003171	1.3	2.28	2.23	2.62	3.09	2.53		
	HEMBA1003175	1.54	2.63	4.2	3.54	4.52	4.11		
20	HEMBA1003179	4.66	5.95	37.4	36.91	43.86	45.13		
	HEMBA1003186	2.58	3.17	7.13	6.71	6.71	5.78		
	HEMBA1003196	3.04	3.79	7.33	6.95	8.31	5.18		
25	HEMBA1003197	0.46	1.51	2.86	1.85	3.97	1.09		
	HEMBA1003199	1.26	1	2.32	1.66	3.22	2.47		
	HEMBA1003202	2.86	3.49	5.69	9.44	10.48	11.14	**	+
	HEMBA1003204	1.67	2.46	3.35	4.99	4.72	4.81	**	+
30	HEMBA1003210	6.48	7.36	11.66	12.02	12.1	14.78		
	HEMBA1003212	1.4	2.87	5.52	7.58	8	5.7		
	HEMBA1003218	1.2	1.26	1.71	1.24	4.35	1.36		
35	HEMBA1003220	34.65	32.6	73.43	78.35	79.82	83.89		
	HEMBA1003222	2.37	3.03	3.41	3.04	6.13	4.29		
	HEMBA1003225	1.95	2.07	3.34	1.59	3.45	2.05		
	HEMBA1003229	2.37	1.91	2.4	5.62	5.1	4.9	**	+
40	HEMBA1003230	7.83	7.14	12.08	11.08	11.44	10.09		
	HEMBA1003235	0.91	1.33	4.32	4.98	5.25	5.44	*	+
	HEMBA1003236	5.54	5.43	10.62	11.5	15.4	13.97	*	+
	HEMBA1003250	1.41	1.4	2.68	1.76	2.98	2.42		
45	HEMBA1003252	4.96	7.17	16.59	17.06	18.68	14.37		
	HEMBA1003257	2.7	3.33	7.33	8.25	8.83	6.78		
	HEMBA1003268	0.95	0.44	1.92	1.92	3.36	1.81		
50	HEMBA1003273	1.4	1.38	2.96	2.5	3.37	5.1		
	HEMBA1003276	1.13	1.99	3.18	4.21	4.42	3.98	*	+
	HEMBA1003277	0.95	0.83	1.85	0.56	1.63	1.34		
	HEMBA1003278	1.07	1.18	3.49	1	4.56	2.32		
55	HEMBA1003280	2.37	2.6	4.59	3.08	4.91	4.2		

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	HEMBA1002986	2.72	4.88	6.67	14.7	14.62	13	**	+
	HEMBA1002988	1.77	2.36	4.25	4.2	5.67	3.46		
5	HEMBA1002992	8.73	11.38	68.65	83.81	96.4	94.02	*	+
	HEMBA1002995	6.13	6.97	11.94	8.64	11.47	14.09		
	HEMBA1002997	5.77	6.33	9.6	12.88	10.65	8.75		
	HEMBA1002999	1.36	2.77	2.84	2.48	3.92	3.31		
10	HEMBA1003004	0.78	1.39	1.96	1.86	3.32	1.37		
	HEMBA1003006	2.03	1.84	4.26	5.44	8.08	5.87	*	+
	HEMBA1003008	1.58	1.26	2.83	3.4	5.4	2.28		
15	HEMBA1003021	1.72	2.09	5.98	8.49	8.58	6.67	*	+
	HEMBA1003027	1.79	1.73	4.47	2.11	6.17	3.72		
	HEMBA1003029	16.39	17.36	46.06	37.07	42.91	45.58		
	HEMBA1003031	33.04	32.41	50.08	48.18	24.56	40.3		
20	HEMBA1003032	3.42	6.52	7.98	8.81	6.53	9.45		
	HEMBA1003033	2.36	4.11	6.85	7.85	8.94	9	*	+
	HEMBA1003034	2.43	3.17	7.63	8.24	7.47	8.23		
25	HEMBA1003035	1.24	2	2.59	2.88	3.46	1.93		
	HEMBA1003037	1.74	2.09	6.21	4.13	7.36	3.43		
	HEMBA1003041	3.4	4.14	8.51	10.28	10.48	9		
	HEMBA1003046	11.44	11.53	28.31	33.77	21.19	36.32		
30	HEMBA1003047	2.02	2.35	5.11	4.57	5.41	4.03		
	HEMBA1003048	1.8	2.96	3.76	7.97	7.47	9.3	**	+
	HEMBA1003064	3.7	4.12	15.74	15.78	25.09	19.36		
	HEMBA1003067	1.92	2.31	7.09	4.56	7.96	3.23		
35	HEMBA1003071	5.24	5	8.74	11.32	10.02	8.7		
	HEMBA1003072	2.81	3.22	5.7	4.43	3.65	5.17		
	HEMBA1003076	20.6	21.34	31.6	41.86	28.3	32.74		
40	HEMBA1003077	1.41	1.58	4.37	1.68	4.03	2.27		
	HEMBA1003078	2.02	1.92	2.4	3.14	4.9	3.38	*	+
	HEMBA1003079	2.72	2.66	6.42	5.88	7.13	4.48		
	HEMBA1003083	1.56	2.11	3.94	4.42	6.23	4.59		
45	HEMBA1003086	2.5	2.72	5.27	4.09	5.78	4.68		
	HEMBA1003090	5.14	4.79	13.3	11.57	12.88	12.73		
	HEMBA1003094	0.82	1.67	2.94	2.51	3.22	2.14		
50	HEMBA1003096	8.6	8.76	15.55	10.1	13.7	10.97		
	HEMBA1003098	3.88	5.66	7.38	9.42	7.11	8.4		
	HEMBA1003101	4.73	5.48	7.29	9.04	6.59	6.36		
	HEMBA1003109	2.88	3.42	4.72	5.73	6.93	7.22	*	+
55	HEMBA1003114	2.87	4.67	5.67	6.47	7.94	5.69		

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	HEMBA1002770	5.89	5.83	12.41	14.24	22.53	15.53	*	+
	HEMBA1002777	1.6	1.9	2.58	2.29	4.74	3.76		
5	HEMBA1002779	10.92	7.6	16	17.39	19.81	19.36	*	+
	HEMBA1002780	2.6	2.77	6.82	6.43	6.89	6.35		
	HEMBA1002790	3.14	2.52	10.6	7.26	8.67	9.25		
	HEMBA1002794	1.52	2.28	5.49	3.68	6.8	4.45		
10	HEMBA1002798	1.33	1.59	3.61	2.77	5.12	4		
	HEMBA1002801	2.13	2.25	3.64	3.12	6.93	5.24		
	HEMBA1002810	4.56	3.99	7.85	10.27	17.31	11.1	*	+
15	HEMBA1002816	2.24	1.97	2.88	5.34	5.05	4.8	**	+
	HEMBA1002818	24.6	23.26	95.11	130.84	121.74	135.78	*	+
	HEMBA1002820	1.95	2.63	6.41	6.96	6.99	6.04		
	HEMBA1002826	1.96	1.48	2.99	3.21	4.84	3.59		
20	HEMBA1002833	8.71	7.46	19.84	20.18	21.16	20.04		
	HEMBA1002850	1.16	1.94	3.67	3.87	4.96	5.11	*	+
	HEMBA1002862	9.06	9.31	17.9	20.11	25.3	13.43		
25	HEMBA1002863	2.47	2.93	5.28	6.16	8.44	6.52	*	+
	HEMBA1002867	1.51	1.17	2.4	2.3	3.28	1.87		
	HEMBA1002876	3.9	3.54	5.48	5.61	5.78	6.48		
	HEMBA1002886	1.28	1.56	2.45	1.83	3.13	2.71		
30	HEMBA1002896	5.82	3.82	9.38	7.22	11.23	8.51		
	HEMBA1002913	2.37	2.22	4.56	4.19	4.28	3.11		
	HEMBA1002921	0.97	0.81	2.36	1.82	2.41	1.41		
	HEMBA1002924	1.07	1.2	2.86	2.11	4.41	3.27		
35	HEMBA1002934	6.01	5.17	10.48	9.93	15.27	13.16		
	HEMBA1002935	4.27	2.55	6.59	7.1	5.34	7.14		
	HEMBA1002937	4.61	5.71	9.4	10.82	8.36	7.36		
40	HEMBA1002939	2.21	2.92	5.39	5.51	5.7	3.26		
	HEMBA1002944	1.45	1.97	4.66	3.1	5.68	3.21		
	HEMBA1002951	5.88	7.88	10.99	6.04	12.17	5.67		
	HEMBA1002954	2.4	4.57	6.12	6.09	7.78	4.78		
45	HEMBA1002962	3.93	6.02	9.14	13.42	15.92	12.44	*	+
	HEMBA1002968	1.22	1.71	4.32	5.34	4.07	5.3		
	HEMBA1002970	1.13	1.13	3.14	2.5	3.72	3.13		
50	HEMBA1002971	0.96	2.02	2.75	2.02	3.71	2.43		
	HEMBA1002973	1.68	3.36	7.84	6.19	10.81	6.31		
	HEMBA1002978	2.09	3.81	4.35	5.49	5.3	4.22		
	HEMBA1002981	1.82	2.51	4.01	9.33	7.48	6.53	**	+
55	HEMBA1002985	0.83	1.92	4.91	4.74	5.59	4.13		

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	HEMBA1002609	4.47	6.27	51	68.51	85.44	66.33	*	+
	HEMBA1002617	6.31	4.76	7.99	7.25	6.84	6.48		
5	HEMBA1002619	3.33	4.8	-5.99	7.86	6.14	7.27		
	HEMBA1002621	1.21	2.94	3.09	3.24	4.17	2.03		
	HEMBA1002624	4.6	5.19	19.48	22.04	24.39	26.87	*	+
	HEMBA1002628	3.37	3.64	6.41	6.08	6.11	4.1		
10	HEMBA1002629	2.71	2.24	4.66	3.77	7.98	4.48		
	HEMBA1002632	1.39	2.23	5.16	4.29	5.49	4.58		
	HEMBA1002645	1.77	1.98	6.43	4.68	6.91	5.37		
15	HEMBA1002651	1.87	2.73	4.73	4.68	4.83	3.74		
	HEMBA1002652	3.38	5.27	6.21	6.09	8.66	7.92		
	HEMBA1002659	2.84	3.86	4.8	6.32	8.18	9.6	*	+
	HEMBA1002661	3	2.71	6.19	4.41	6.93	4.93		
20	HEMBA1002666	1.74	2.47	4.21	2.95	4.25	1.41		
	HEMBA1002667	1.39	2.25	3.91	2.79	5.24	1.94		
	HEMBA1002673	16.08	19.36	30.31	32.54	35.18	29.96		
25	HEMBA1002678	2.11	2.33	7.44	5.39	5.98	4.22		
	HEMBA1002679	1.23	2.33	5.25	3.7	7.48	3.81		
	HEMBA1002688	1.74	2.98	8.3	8.33	11.41	7.86		
	HEMBA1002696	1.7	2.79	2.92	3.48	6.13	3.32		
30	HEMBA1002703	2.95	3.88	10.15	8.35	9.73	9.21		
	HEMBA1002706	4.97	4.24	8.99	5.07	7.16	5.54		
	HEMBA1002712	2.39	3.94	8.67	8.4	10.9	10.57		
35	HEMBA1002715	7.92	9.81	49.65	79.65	93.63	79.61	*	+
	HEMBA1002716	3.93	4.26	5.53	4.63	5.02	4.53		
	HEMBA1002718	11.79	12.87	17.77	24.16	18.07	24.3	*	+
	HEMBA1002728	2.37	3.1	5.01	5.52	5.94	4.42		
40	HEMBA1002730	1.13	2.48	5.86	3.71	6.19	4.61		
	HEMBA1002734	2.89	3.54	8.82	8.6	10.7	10.59		
	HEMBA1002742	1.94	2.06	3.96	1.86	4.27	2.74		
	HEMBA1002746	1.2	2.86	4.61	2.83	4.43	2.94		
45	HEMBA1002748	2.19	1.75	4.01	5.36	5.98	3.92		
	HEMBA1002750	1.99	2.46	3.45	6.74	6.39	6.27	**	+
	HEMBA1002755	1.85	3.1	5.31	5.96	6.62	5.16		
50	HEMBA1002759	1.93	3.12	7.98	4.65	7.92	7.08		
	HEMBA1002763	9.62	12.05	74.52	68.84	88.82	77.22		
	HEMBA1002767	4.48	5.85	5.8	8.88	6	6.13		
	HEMBA1002768	2.99	3.76	6.2	3.46	8.3	3.04		
55	HEMBA1002769	1.47	2.35	2.82	3.46	5.21	3.49		

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	HEMBA1002462	5.18	3.83	9.52	11.92	9.29	8.58		
	HEMBA1002465	0.93	1.96	2.26	1.46	3.13	1.27		
5	HEMBA1002469	6.88	7.27	45.87	49.02	75.59	67.74	*	+
	HEMBA1002475	1.54	2.35	8.01	4.88	7.87	7.66		
	HEMBA1002477	1.75	1.59	5.25	3.19	6.99	3.55		
	HEMBA1002480	4.46	3.98	8.49	8.76	13	10.54		
10	HEMBA1002481	1.9	2.02	4.22	2.71	6.25	4.7		
	HEMBA1002486	3.62	3.98	10.08	9.98	8.04	9.75		
	HEMBA1002490	2.02	3.08	5.7	8.76	9.64	7.65	*	+
15	HEMBA1002495	2.37	2.29	3.78	3.92	4.79	4.08		
	HEMBA1002498	0.95	2.14	2.97	1.83	5.09	2.14		
	HEMBA1002501	2.96	4.73	14.13	19.98	23.55	17.54	*	+
	HEMBA1002503	1.7	2.52	5.11	4.68	7.06	2.97		
20	HEMBA1002504	1.95	2.19	5.99	6.68	7.09	4.65		
	HEMBA1002508	1.48	2.59	5.99	7.8	7.47	5.65		
	HEMBA1002513	1.31	1.7	4.85	3.91	7.67	3.02		
25	HEMBA1002515	1.17	1.82	3.04	2.67	5.1	2.89		
	HEMBA1002524	1.67	2.09	2.53	4.44	4.49	3.82	**	+
	HEMBA1002538	4.68	4.14	7.39	9.31	8.91	7.86	*	+
	HEMBA1002542	3.31	3.27	6.77	10.11	9.3	7.74	*	+
30	HEMBA1002544	1.42	2.24	3.33	2.69	6.59	3.24		
	HEMBA1002546	31.01	31.64	56.69	95.52	83.15	72.77	*	+
	HEMBA1002547	3.13	3.22	8.44	20.11	20.37	17.21	**	+
35	HEMBA1002550	5.46	3.86	10.87	10.85	11.2	10.23		
	HEMBA1002551	2.15	3.09	5.8	3.7	5.08	3.08		
	HEMBA1002552	2.21	2.06	8.39	5.66	6.55	5.68		
	HEMBA1002555	1.54	1.78	4.56	2.27	4.4	2.97		
40	HEMBA1002558	2.74	3.26	7.02	8.08	7.47	8.27		
	HEMBA1002561	1.01	1.58	5.26	4.42	5.08	3.87		
	HEMBA1002562	0.59	0.83	2.34	3.29	3.29	2.36		
	HEMBA1002568	1.71	1.16	3.09	3.06	3.26	3.6		
45	HEMBA1002569	3.8	4.67	10.32	7.29	8.59	5.14		
	HEMBA1002570	5.22	4.72	9.84	6.07	10.29	12.99		
	HEMBA1002574	24.62	22.75	26.01	44.47	30.74	40.85	*	+
50	HEMBA1002583	4.07	4.52	8.07	6.64	6.43	8.47		
	HEMBA1002587	9.78	10.9	19.23	24.67	18.08	20.4		
	HEMBA1002590	2.51	2.58	7.47	5.35	5.6	4.05		
	HEMBA1002592	2.51	3.03	6.4	6.34	7.84	5.1		
55	HEMBA1002595	1.66	2.13	3.1	4.12	4.25	2.68		

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	HEMBA1002319	1.97	2.96	4.46	8.32	9.58	8	**	+
	HEMBA1002320	1.99	1.76	3.22	2.51	4.45	1.76		
5	HEMBA1002321	1.22	2.04	-2.84	2.71	4.44	2.77		
	HEMBA1002328	2.44	3.04	4.89	2.43	5.73	2.94		
	HEMBA1002333	3.88	4.27	7.14	7.37	10.84	7.5		
	HEMBA1002337	3.02	3.62	5.5	5.98	7.49	7.48	*	+
10	HEMBA1002339	15.86	13.92	111.66	135.44	169.95	156.76	*	+
	HEMBA1002341	0.8	2.08	3.22	2.71	4.34	2.35		
	HEMBA1002348	2.84	2.78	7.14	3.69	6.73	4.49		
15	HEMBA1002349	1.28	1.44	3.59	2.24	4.64	2.59		
	HEMBA1002353	1.83	3.04	4.03	4.61	7.72	5.68		
	HEMBA1002356	6.05	6.96	17.53	14.27	16.02	16.1		
	HEMBA1002357	114.85	156.08	306.32	300.67	286.5	328.19		
20	HEMBA1002360	7.18	8.32	8.29	14.57	14.46	13.78	**	+
	HEMBA1002363	2.79	3.35	4.84	7.02	8.02	8.72	**	+
	HEMBA1002365	1.7	2.7	2.7	1.63	3.12	2.67		
25	HEMBA1002370	1.43	1.78	2.37	1.53	4.2	1.9		
	HEMBA1002374	4.55	4.53	7.79	8.33	10.27	9.11	*	+
	HEMBA1002376	46.59	33.18	118.8	101.1	189.18	114.36		
	HEMBA1002377	18.02	20.98	25.61	32.58	34.44	32.19	**	+
30	HEMBA1002380	5.68	6.36	16.28	17.43	21.85	18.83		
	HEMBA1002381	1.52	1.8	4.16	4.12	7.16	4.94		
	HEMBA1002384	1.79	3.09	3.69	5.67	4.27	5.71	*	+
	HEMBA1002389	1.93	2.93	2.88	3.63	5.31	4.7	*	+
35	HEMBA1002396	21.16	20.01	36.93	14.29	14.94	19.1		
	HEMBA1002402	125.09	124.52	168.42	100.85	107.79	164.62		
	HEMBA1002417	1.41	1.07	4.27	2.17	3.19	2.6		
40	HEMBA1002419	1.42	2.38	3.8	1.81	4.59	2.41		
	HEMBA1002420	9.55	11.97	14.11	16.34	18.28	16.16	*	+
	HEMBA1002421	7.47	10.35	12.5	8.97	10.24	9.6		
	HEMBA1002423	2.89	1.3	4.28	3.35	4.29	3.82		
45	HEMBA1002424	11.91	10.05	25.13	9.11	9.2	10.43		
	HEMBA1002426	3.42	3.69	5.56	7.57	8.24	5.6	*	+
	HEMBA1002430	0.39	1.41	2.51	1.7	4.19	2.85		
	HEMBA1002439	1.59	1.94	4.17	2.69	4.46	6.8		
50	HEMBA1002441	31.85	29.77	27.79	48.74	52.7	34.64		
	HEMBA1002454	0.62	1.48	2.27	1.76	2.43	1.55		
	HEMBA1002458	3.17	5	8.09	10.24	10.55	11.71	*	+
55	HEMBA1002460	2.14	1.59	3.89	5.16	4.63	5.01	*	+

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	HEMBA1002166	21.88	18.32	39.58	57.35	49.05	46.09	*	+
	HEMBA1002167	0.89	2.89	3.89	4.96	5.45	3.98		
5	HEMBA1002173	3.24	3.83	6.22	7.97	7.11	6.28		
	HEMBA1002177	1.31	1.78	3.31	5.68	4.97	2.98		
	HEMBA1002178	6.91	10.17	14.77	23.33	23.58	17.49	*	+
10	HEMBA1002179	53.56	46.86	94.4	58.33	85.22	54.47		
	HEMBA1002185	2.75	4.07	13.4	11.73	16.23	14.56		
	HEMBA1002188	5.76	7.57	10.27	11.86	12.9	9.8		
	HEMBA1002189	1.98	2.85	4.96	5.23	4.63	4.71		
15	HEMBA1002191	0.67	2.16	4.96	3.47	5.44	2.81		
	HEMBA1002192	2.98	2.83	4.91	7.53	8.35	4.57		
	HEMBA1002195	2.96	3.27	6.6	10.35	10.11	7.27	*	+
	HEMBA1002196	3.34	4.33	8.55	8.62	8.85	8.39		
20	HEMBA1002199	1.33	1.86	4.9	4.62	5.71	3.52		
	HEMBA1002204	1.31	1.97	4.08	5.48	11.37	3.73		
	HEMBA1002208	24.58	26.61	45.85	49.77	25.48	39.6		
25	HEMBA1002212	3.73	5.95	9.01	8.9	11.85	17.18		
	HEMBA1002215	1.95	2.63	4.27	5.1	3.54	3.78		
	HEMBA1002217	15.61	16.71	59.91	78.46	82.88	80.94	*	+
	HEMBA1002220	1.11	2.07	4.1	3.58	3.39	2.33		
30	HEMBA1002226	2.17	3.13	9.18	10.47	12.61	9.58		
	HEMBA1002227	39.9	47.13	92.5	109.42	65.74	71.79		
	HEMBA1002229	4.5	4.77	13.39	11.16	13.55	12.49		
35	HEMBA1002237	1.73	3.22	4.08	3.71	5.64	3.41		
	HEMBA1002239	9.36	13.83	72.18	100.62	109.3	113.84	*	+
	HEMBA1002241	7	7.54	38.36	64.27	68.93	68.72	**	+
	HEMBA1002253	1.11	2.44	3.33	2.3	4.42	2.68		
40	HEMBA1002257	1.83	2.65	4.11	3.18	3.6	1.74		
	HEMBA1002259	1.12	2.17	2.69	3.12	3.6	2.67		
	HEMBA1002262	6.95	7.37	19.16	14.43	14.78	17.04		
	HEMBA1002265	1.35	1.63	3.7	3.75	6.23	2.43		
45	HEMBA1002267	16.87	20.81	22.76	32.99	16.96	27.5		
	HEMBA1002270	3.73	4.79	7.49	8.18	13.43	8.7		
	HEMBA1002286	1.03	1.86	5.42	2.85	5.53	0.98		
50	HEMBA1002290	4.73	3.7	7.52	5.16	8.88	4.38		
	HEMBA1002302	6.12	9.63	45.04	45.66	49.69	45.23		
	HEMBA1002304	3.28	3.42	6.88	5.57	6.97	4.34		
	HEMBA1002307	45.71	53.69	92.31	71.87	55.79	61.03		
55	HEMBA1002316	2.16	3.29	4.63	3.04	5.32	2.41		

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	HEMBA1002037	4.47	4.34	6.69	4.5	6.47	7.94		
	HEMBA1002038	4.12	3.13	7.74	6.36	8.8	5.42		
5	HEMBA1002039	2.46	3.43	7.03	6.74	8.99	6.37		
	HEMBA1002042	5.52	5.55	9.8	10.55	13.01	9.94		
	HEMBA1002043	3.81	3.79	11.32	12.53	12.64	13.95		
	HEMBA1002048	2.76	2.31	3.81	2.03	4.32	3.44		
10	HEMBA1002049	1.72	2.35	5.55	4.43	5.2	5.2		
	HEMBA1002053	7.33	6.91	14.52	11.09	15.25	13.63		
	HEMBA1002055	9.81	8.76	10.65	18.44	13.58	17.78	*	+
15	HEMBA1002056	2.24	2.62	4.26	5.46	8.06	5.67	*	+
	HEMBA1002061	2.24	2.51	4.58	4.17	5.34	4.58		
	HEMBA1002080	46.55	49.5	54.6	91.78	122.41	83.23	*	+
	HEMBA1002084	0.71	1.43	2.36	3.25	3.66	2.64	*	+
20	HEMBA1002085	0.97	1.47	2.87	3.45	4.52	3.74	*	+
	HEMBA1002092	1.79	1.56	3.94	4.01	5.53	4.15		
	HEMBA1002098	1.51	1.82	4.12	3.2	5.11	2.83		
25	HEMBA1002100	9.07	8.18	22.37	25.95	23.04	29.67		
	HEMBA1002101	18.26	17.64	27.49	23.44	27.16	23.74		
	HEMBA1002102	2.65	1.98	5.99	4.58	7.06	6.99		
	HEMBA1002105	6.79	6.2	22.13	24.47	40.8	25.31		
30	HEMBA1002107	57.97	37.86	84.35	155.21	136.27	136.87	**	+
	HEMBA1002113	6.77	4.75	17.24	12.05	12.83	14.78		
	HEMBA1002119	3.85	3.28	24.05	24.29	30.66	27.04		
35	HEMBA1002125	7.03	6.73	10.43	11.38	7.34	11.85		
	HEMBA1002131	9.71	9.72	20.58	14.97	19.12	13.71		
	HEMBA1002133	3.67	3.52	6.32	6.97	9.51	9.46	*	+
	HEMBA1002139	0.75	1.07	3.35	1.52	4.37	1.47		
40	HEMBA1002141	1.67	1.36	3.23	2.99	3.92	2.14		
	HEMBA1002144	2.33	2.44	6.11	5.28	5.68	6.57		
	HEMBA1002147	8.84	9.55	17.93	36.22	21.92	21.88		
45	HEMBA1002150	38.34	38.68	51.42	19.74	26.62	15.85	*	-
	HEMBA1002151	3.76	3.36	9.95	11.15	14.23	9.84		
	HEMBA1002153	0.57	1.74	3.36	3.39	5.64	2.96		
	HEMBA1002156	0.8	1.74	2.33	1.94	4.28	1.26		
50	HEMBA1002160	2.16	3.17	5.7	6.08	7.26	6.18		
	HEMBA1002161	2.13	2.9	6.99	13.53	11.79	9.76	*	+
	HEMBA1002162	2.65	2.17	7.76	5.61	6.27	7.81		
	HEMBA1002163	12.02	12.04	19.93	34.48	19.96	27.11		
55	HEMBA1002164	6.58	10.55	59.92	71.46	78.61	67.82		

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	HEMBA1001879	1.89	2.57	5.58	5.99	7.24	5.72		
	HEMBA1001884	6.21	6.49	17.14	11.31	12.71	10.87		
5	HEMBA1001886	2.12	2.21	4.38	4.57	5.67	4.23		
	HEMBA1001888	2	2.12	6.6	7.41	10.17	9.46	*	+
	HEMBA1001890	4.03	3.67	7.6	6.8	7.01	4.4		
10	HEMBA1001896	1.34	1.61	2.62	2.27	4.12	3.01		
	HEMBA1001899	33.43	39.48	61.77	106.52	41.01	101.91		
	HEMBA1001904	76.64	122.45	233.99	299.33	174.47	322.82		
	HEMBA1001910	1.4	1.93	3.23	2.53	6.35	2.97		
15	HEMBA1001911	8.36	8.75	10.86	21.15	16.9	13.23	*	+
	HEMBA1001912	8.92	7.97	33.97	57	51.9	48.59	*	+
	HEMBA1001913	4.89	6.19	17.29	18.56	14.16	16.85		
20	HEMBA1001915	1.35	2.61	4.49	3.3	5.63	2.46		
	HEMBA1001918	15.23	13.29	21.07	17.07	14.31	12.13		
	HEMBA1001921	4	3.5	4.38	5.2	5.35	4.86	*	+
	HEMBA1001931	1.19	1.95	2.53	2.14	5.17	2.19		
25	HEMBA1001939	1.92	1.77	4.72	1.97	5.21	2.57		
	HEMBA1001940	2.61	2.99	7.14	3.51	5.86	3.24		
	HEMBA1001942	1.18	1.88	3.71	2.33	5.14	1.56		
30	HEMBA1001944	4.35	5.83	42.16	51.42	66.43	59.75	*	+
	HEMBA1001945	0.98	2.3	2.95	2.98	3.4	2.21		
	HEMBA1001950	2.56	2.84	7.87	5.72	5.23	3.68		
	HEMBA1001951	10.37	11.26	15.33	24.16	18.26	22.94	*	+
35	HEMBA1001958	1.04	1.28	2.58	3.1	4.83	2.54		
	HEMBA1001960	6.87	6.28	13.93	10.02	12.99	12.47		
	HEMBA1001962	1.01	1.08	4.19	1.58	4.24	1.67		
	HEMBA1001964	1.39	3.45	4.13	2.54	4.45	3.39		
40	HEMBA1001967	6.06	5.65	9.33	14.45	10.5	13.18	*	+
	HEMBA1001979	0.7	2.67	3.31	2.04	3.46	2.4		
	HEMBA1001987	1.96	3.92	7.99	6.19	8.35	7.22		
45	HEMBA1001991	1.61	3.59	9.06	5.06	8.7	7.44		
	HEMBA1002003	4.86	4.71	14.56	15.86	16.03	22.9		
	HEMBA1002005	2.62	3.39	7.82	4.16	7.48	4.76		
	HEMBA1002008	2.64	3.51	7.78	6.07	10.15	6.37		
50	HEMBA1002018	1.86	2.37	4.23	3.32	5.47	3.87		
	HEMBA1002022	0.52	2.3	2.5	2.83	3.53	2.82		
	HEMBA1002029	43.82	40.22	73.75	89.27	96.12	122.81	*	+
	HEMBA1002030	2.23	2.88	4.32	3.88	4.26	4.67		
55	HEMBA1002035	1.69	1.75	3.82	5.43	5.14	3.75		

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	HEMBA1001723	3.43	3	10.19	9.09	12.53	9.64		
	HEMBA1001731	1.3	1.36	3.27	1.58	4.8	2.29		
5	HEMBA1001734	2.37	2.38	4.28	4.78	6.06	4.27		
	HEMBA1001736	2.3	2.12	2.87	3.5	4.69	5.33	*	+
	HEMBA1001741	1.69	2.19	3.5	4.02	4.74	2.9		
10	HEMBA1001744	0.86	0.94	2.81	2	3.64	2.64		
	HEMBA1001745	0.95	1.56	2.3	2.53	5.28	2.58		
	HEMBA1001746	4.02	3.91	8.66	9.21	12.06	7.01		
	HEMBA1001761	2.2	2.01	4.69	2.58	3.95	3.37		
15	HEMBA1001762	1.41	1.6	3.57	1.93	4.38	2.03		
	HEMBA1001781	1.56	1.5	4.17	2.15	5.87	3.25		
	HEMBA1001784	1.36	1.39	4.5	4.02	3.58	3.75		
	HEMBA1001791	2.16	1.74	6.97	6.04	7.62	5.68		
20	HEMBA1001794	2.15	4.31	12.57	13.54	13.65	12.96		
	HEMBA1001800	5.61	9.63	60.44	84.85	100.03	76.26	*	+
	HEMBA1001803	2.84	4.25	5.36	4.27	7.02	3.67		
25	HEMBA1001804	6.2	8.13	20.95	29.84	26.86	24.39	*	+
	HEMBA1001808	1.61	1.6	3.87	3.71	3.67	2.89		
	HEMBA1001809	8.07	6.27	10.64	14.33	20.56	16.63	*	+
	HEMBA1001811	8.32	7.83	16.8	22.75	21.75	17.6	*	+
30	HEMBA1001815	1.75	2.67	6.56	5.58	6.33	5.03		
	HEMBA1001816	1.96	2.67	4.47	3.09	4.6	3.04		
	HEMBA1001819	0.98	3.09	6.16	6.19	8.53	6.3		
35	HEMBA1001820	0.93	1.32	2.22	2.36	3.32	1.21		
	HEMBA1001822	1.87	2.06	5.43	6.02	7.7	4.44		
	HEMBA1001824	3.21	4.62	14.88	12.81	16.29	12.34		
	HEMBA1001835	1.04	1.05	3.05	3.72	5.21	3.14		
40	HEMBA1001844	7.88	6.55	18.04	17.77	21.36	13.19		
	HEMBA1001847	0.93	1.8	5.21	1.96	5.18	3.06		
	HEMBA1001849	2.32	2.77	7.58	6.65	8.19	7.62		
	HEMBA1001850	2.51	2.71	8.43	8.76	8.88	7.89		
45	HEMBA1001861	0.95	2.04	1.73	2.64	3.93	2.01		
	HEMBA1001862	138.58	133.42	191.61	266.65	221.43	227.58	*	+
	HEMBA1001864	1.31	1.16	2.44	4.79	2.88	2.59		
50	HEMBA1001866	1.49	2.39	7.45	7.67	7.07	4.61		
	HEMBA1001869	7.55	6.84	10.82	10.31	7.69	9.02		
	HEMBA1001871	29.48	30.98	54.77	63.07	62.43	66.59	*	+
	HEMBA1001876	0.96	1.27	4.42	2.08	4.57	2.26		
55	HEMBA1001878	2.23	3.34	5.7	6.83	8.18	5.5		

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	HEMBA1001569	11.15	11.91	26.6	30.2	31.14	32.61	*	+
	HEMBA1001570	3.25	4.61	10.2	9.19	10.25	9.53		
5	HEMBA1001579	3.63	4.4	-7.77	9.26	7.4	9.93		
	HEMBA1001581	2.79	3.33	10.95	8.81	12.09	9.08		
	HEMBA1001582	3.22	3.18	6.68	6.35	6.84	4.03		
10	HEMBA1001585	2.7	3.07	4.52	4.4	5.6	3.5		
	HEMBA1001589	1.82	2.31	3.63	4.39	6.19	3.78		
	HEMBA1001595	13.06	15.57	19.7	13.25	13.29	14.02		
	HEMBA1001604	1.96	2.67	3.64	3.76	6.53	2.82		
15	HEMBA1001608	5.58	7.09	16.17	14.14	16.46	14.43		
	HEMBA1001615	113.28	90.33	205.41	240.97	118.65	165.59		
	HEMBA1001620	3.71	5.56	10.54	12.22	12.24	11.46		
20	HEMBA1001621	0.76	2.13	3.42	1.76	3.44	2.97		
	HEMBA1001635	2.32	2.13	3.41	3.55	4.9	2.85		
	HEMBA1001636	1.9	1.93	4.01	3.34	5.33	2.97		
	HEMBA1001640	3.07	3.31	13.65	10.96	15.01	10.74		
25	HEMBA1001647	8.92	8.44	57.38	88.92	112.42	87.46	*	+
	HEMBA1001651	2.53	3.54	7.85	6.62	9.07	8.73		
	HEMBA1001655	2.09	2.66	4.78	3.35	6.75	4.09		
	HEMBA1001658	4.33	4.5	9.27	7.26	11.15	8.6		
30	HEMBA1001661	0.75	1.78	2.8	1.98	3.22	1.77		
	HEMBA1001665	1.52	1.85	3.47	2.63	6.63	1.73		
	HEMBA1001670	5.32	6.54	8.82	12.45	15.21	12.42	**	+
35	HEMBA1001672	2.49	3.06	5.9	4.28	7.62	3.39		
	HEMBA1001673	8.23	10.76	13.22	20.04	19.39	15.65	*	+
	HEMBA1001675	2.4	2.01	2.53	3.21	5.79	3.36		
	HEMBA1001676	54.19	46.09	107.65	245.72	212.81	275.65	**	+
40	HEMBA1001678	9.46	10.2	21.87	23.65	19.51	27.88		
	HEMBA1001680	4.58	4.89	12.32	9.39	10.95	11.65		
	HEMBA1001681	1.71	2.44	5.75	6.25	9.11	6.36		
45	HEMBA1001684	1.89	2.74	6.26	4.32	7.57	6.98		
	HEMBA1001695	1.48	2.08	3.42	2.3	4.76	3.15		
	HEMBA1001702	1.54	2.96	3.55	2.36	7.57	3.09		
	HEMBA1001709	1.23	1.8	3.51	3.21	4.87	3.5		
50	HEMBA1001711	1.29	1.98	2.83	2.99	2.45	3.18		
	HEMBA1001712	0.92	1.55	2.56	2.13	3.02	2.24		
	HEMBA1001714	10.37	10.82	19.06	23.54	22	23.8	*	+
	HEMBA1001717	79.4	71.16	124.25	152.62	195.81	173.65	*	+
55	HEMBA1001718	1.95	2.12	7.32	5.99	6.59	5.26		

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	HEMBA1001407	1.13	1.78	3.73	6.39	6.64	4.44	*	+
	HEMBA1001411	1.44	2.81	4.47	7.2	6.35	6.04	*	+
5	HEMBA1001413	1.84	1.53	-3.31	3.61	3.75	3.76		
	HEMBA1001414	1.47	2.34	5.3	5.33	7.22	5.47		
	HEMBA1001415	1.91	2.36	4.97	4.4	5.86	4.29		
10	HEMBA1001416	4.73	4.85	9.54	8.87	11.06	9.4		
	HEMBA1001432	1.23	1.27	4.43	4.27	6.64	3.59		
	HEMBA1001433	1.96	2.93	4.55	4.33	8.66	3.64		
	HEMBA1001435	2.17	2.27	6.39	7.02	10.35	5.88		
15	HEMBA1001442	0.99	0.68	2.02	2.36	3.12	1.81		
	HEMBA1001446	1.87	1.84	5.82	9.71	8.93	11.01	**	+
	HEMBA1001450	2.35	2.32	11.22	8.61	10.08	6.34		
20	HEMBA1001454	3.08	4.25	9.69	13.64	10.73	11.82	*	+
	HEMBA1001455	2.28	2.7	3.11	2.69	5.54	2.8		
	HEMBA1001459	2.74	3.37	6.03	5.07	7.38	5.52		
	HEMBA1001461	3.34	4.47	6.96	6.8	9.85	7.47		
25	HEMBA1001462	1.07	1.47	2.79	2.67	4.5	2.54		
	HEMBA1001463	1.38	1.61	5.25	4.95	5.46	5.51		
	HEMBA1001469	3.9	4.51	7.32	10.63	9.83	7.76	*	+
	HEMBA1001473	4.56	3.49	8.25	7.52	10.34	6.53		
30	HEMBA1001477	2.14	1.59	4.64	3.41	5.75	2.59		
	HEMBA1001478	2.46	2.8	3.77	2.95	3.73	2.55		
	HEMBA1001480	4.15	6.8	8.96	11.64	11.87	8.48		
35	HEMBA1001483	1.9	1.64	5.71	6.6	8.22	7.23	*	+
	HEMBA1001490	1.45	2.09	3.76	5.16	4.52	4.65	*	+
	HEMBA1001495	56.8	53.41	123.27	193.11	133.65	132.04		
	HEMBA1001497	2.06	1.98	7.47	4.81	8.45	5.85		
40	HEMBA1001510	3.99	4.23	15.22	11.46	13.7	14.56		
	HEMBA1001515	1.45	2.33	4.02	3.73	6.11	3.04		
	HEMBA1001517	1.6	2.21	4.6	5.26	5.4	4.6		
45	HEMBA1001522	1.56	2.72	3.77	3.61	6.37	2.43		
	HEMBA1001526	2.19	2.97	4.97	4.05	4.38	3.59		
	HEMBA1001533	3.19	2.86	6.23	6.83	7.76	4.64		
	HEMBA1001547	7.26	5.37	13.69	5	7.96	6.19		
50	HEMBA1001552	7.12	4.72	17.79	16.12	16.3	16.05		
	HEMBA1001553	41.67	45.48	66.23	57.2	47.01	79.81		
	HEMBA1001557	2.24	2.93	5.15	5.81	8.33	4.59		
	HEMBA1001563	1.69	2.4	4.56	3.66	6.99	4.76		
55	HEMBA1001566	1.42	3.27	8.29	5.94	9.04	5.84		

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	HEMBA1001279	2.54	3.52	13.6	18.68	23.45	18.99	*	+
	HEMBA1001281	16.58	20.99	40.84	47.71	59.04	45.72	*	+
5	HEMBA1001286	3.25	4.71	10.71	11.24	10.65	12.38		
	HEMBA1001289	0.41	1.57	1.64	1.3	3.57	2.41		
	HEMBA1001291	3.52	4.58	9.53	10.91	18.3	18.8	*	+
10	HEMBA1001294	2.01	1.81	4.6	4.04	7.73	5.12		
	HEMBA1001296	3.4	3.52	4.37	3.77	5.94	5.22		
	HEMBA1001297	2.88	3.61	5.51	4.81	6.88	5.38		
	HEMBA1001299	2.49	2.9	6.21	6.45	8.84	7.74	*	+
15	HEMBA1001302	9.42	11.94	15.5	23.25	35.12	25.74	*	+
	HEMBA1001303	1.8	1.99	2.61	3.57	3.8	3.3	**	+
	HEMBA1001306	1.4	1.15	2.85	5.01	4.46	4.82	**	+
	HEMBA1001308	3.43	4.37	16.7	16.31	18.28	21.75		
20	HEMBA1001310	1.93	1.71	4.17	2.38	6.26	3.28		
	HEMBA1001312	10.09	10.35	17.42	20.51	24.71	21.67	*	+
	HEMBA1001319	1.23	1.41	3.85	2.23	4.27	4.01		
25	HEMBA1001322	1.81	2.29	4.17	2.83	4.74	3.78		
	HEMBA1001323	4.04	3.65	8.44	14.68	23.44	18.68	*	+
	HEMBA1001326	8.79	7.35	10.15	12.24	13.62	15.04	*	+
	HEMBA1001327	0.94	1.65	3.18	3.55	5.18	4.56	*	+
30	HEMBA1001330	1.59	2.22	6.96	7.36	9.28	9.64	*	+
	HEMBA1001348	1.68	3.99	3.89	6.33	9.84	7.47	*	+
	HEMBA1001350	5.28	4.16	6.34	7.24	13.17	10.12		
35	HEMBA1001351	15.37	14.99	17.64	37.37	49.52	25.96	*	+
	HEMBA1001352	3.25	3.62	5.97	8.16	13.65	5.75		
	HEMBA1001353	30.24	37.73	49.4	76.74	96.09	96.34	**	+
	HEMBA1001358	13.98	9.73	17.96	30.89	27.69	30.6	**	+
40	HEMBA1001361	1.7	3.24	4.96	4.18	6.08	6.06		
	HEMBA1001364	0.8	1.71	2.4	1.47	4.11	2.95		
	HEMBA1001375	3.45	2.77	5.75	5.71	5.83	6.32		
45	HEMBA1001377	2.81	3.16	7.36	5.37	7.98	7.89		
	HEMBA1001383	0.25	1.64	2.61	1.26	2.47	1.84		
	HEMBA1001387	1.81	2.15	3.66	1.94	5.14	2.47		
	HEMBA1001388	1.52	1.78	5.07	2.01	4.61	3.49		
50	HEMBA1001390	34.61	34.52	66.57	67.03	50	56.4		
	HEMBA1001391	1.65	2.77	4.83	4.32	7.98	3.82		
	HEMBA1001398	1.98	2.87	7.47	7.24	10.42	8.29		
	HEMBA1001405	1.17	2	3.87	2.99	5.3	2.61		
55	HEMBA1001406	2.01	3.27	3.75	5.35	6.62	4.33	*	+

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	HEMBA1001109	4.87	3.77	8.57	11.32	14.48	11.73	*	+
	HEMBA1001114	44.68	41.2	93.35	141.87	145.19	167.76	**	+
5	HEMBA1001121	2.14	2.03	-3.87	2.41	6	3.25		
	HEMBA1001122	9.79	10	14.12	7.73	11.5	22.69		
	HEMBA1001123	2.79	3.28	5.2	5.81	6.02	4.95		
10	HEMBA1001133	0.97	1.69	2.54	2.78	3.84	1.21		
	HEMBA1001137	0.82	1.73	3.65	3.74	3.36	2.54		
	HEMBA1001140	1.23	2.75	2.98	3.62	5.18	4.34	*	+
	HEMBA1001144	4.12	3.41	9.06	14.13	14.12	13.96	**	+
15	HEMBA1001145	47.87	43.87	65.7	98.4	75.15	81.3	*	+
	HEMBA1001158	7.55	9.5	11.62	13.02	7.58	12.5		
	HEMBA1001172	1.44	2.85	4.37	5.32	5.77	5.17	*	+
	HEMBA1001174	0.95	2.06	2.83	3.88	6.31	3.25		
20	HEMBA1001175	6.93	8.56	10.73	14.17	14.5	10.18		
	HEMBA1001182	16.93	19.89	82.44	135.93	145.36	122.22	*	+
	HEMBA1001184	1.41	1.24	2.45	1.85	3.03	1.47		
25	HEMBA1001192	1.72	1.75	4.01	5.65	5.17	3.98		
	HEMBA1001196	2.31	3.63	7.61	9.43	10.51	8.97	*	+
	HEMBA1001197	31.18	35.89	86.14	95.35	83.09	93.59		
	HEMBA1001208	1.83	2.59	3	2.67	5.3	2.61		
30	HEMBA1001213	12.99	16.12	69.9	102.88	119.96	113.72	*	+
	HEMBA1001214	1.39	3.11	4.36	5.14	7.04	4.62		
	HEMBA1001221	1.63	1.62	3.66	2.06	4.19	1.89		
	HEMBA1001225	1.06	2.66	3.53	1.44	3.43	1.52		
35	HEMBA1001226	4.76	4.65	11.94	13.58	15.58	14.92	*	+
	HEMBA1001228	72.4	75.3	102.4	38.23	64.63	78.89		
	HEMBA1001229	18	21.39	82.05	115.91	145.39	128.91	*	+
40	HEMBA1001235	3.58	4.11	6.48	7.31	6.7	10.2		
	HEMBA1001238	2.46	2.49	7.23	4.6	6.94	4.74		
	HEMBA1001242	15.36	14.03	91.45	92.81	94.02	90.34		
	HEMBA1001247	4.41	4.36	12.46	12.48	14.07	15.62		
45	HEMBA1001253	8.79	11.4	61.56	77.17	102.24	94.81	*	+
	HEMBA1001257	1.98	2.71	3.78	3.52	4.29	3.17		
	HEMBA1001261	3.01	3.18	4.56	4.54	3.75	5.59		
50	HEMBA1001262	1.48	3.79	2.81	2.42	4.34	4.59		
	HEMBA1001265	2.76	3.21	6.85	5.5	7.32	5.1		
	HEMBA1001266	3.97	3.17	6.31	7.8	10.5	8.38	*	+
	HEMBA1001269	15.98	10.36	12.79	22.69	24.71	25.21	**	+
55	HEMBA1001272	1.31	2.04	4.3	1.62	5.12	2.07		

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	HEMBA1000968	1.73	1.86	4.7	4.1	4.83	4.66		
	HEMBA1000971	1.75	2.51	2.9	4.18	5.27	5.71	**	+
5	HEMBA1000972	1.45	1.57	3.83	2.63	4.44	3.49		
	HEMBA1000974	1.69	2.69	6.33	7.39	9.35	8.82	*	+
	HEMBA1000975	0.9	1.83	4.17	3.31	5.54	5.12		
	HEMBA1000979	1.45	1.69	3.98	2.55	6.12	3.93		
10	HEMBA1000981	4.21	6.9	9.5	11.75	13.27	14.72	*	+
	HEMBA1000983	1.94	1.45	3.01	3.89	4.53	4.15	*	+
	HEMBA1000985	1.58	0.92	2.75	1.73	3.28	2.79		
15	HEMBA1000986	1.2	1.48	2.47	3.61	4.91	4.26	**	+
	HEMBA1000991	1.56	1.86	3.8	3.11	5.05	5.96		
	HEMBA1001007	0.89	1.08	4.08	1.84	3.89	2.71		
	HEMBA1001008	3.64	3.41	5.86	3.89	7.89	4.95		
20	HEMBA1001009	0.89	1.3	3.07	1.58	3.83	1.81		
	HEMBA1001014	3.54	4.39	9.91	11.82	15.38	14.12	*	+
	HEMBA1001017	4.21	2.82	5.6	6.04	5.41	8.55		
25	HEMBA1001019	1.92	2.81	3.97	8.71	7.74	8.29	**	+
	HEMBA1001020	1.23	2.71	2.3	2.84	5.05	3.6		
	HEMBA1001021	1.07	1.62	2.89	3.13	5.24	2.63		
	HEMBA1001022	2.29	2.25	4.35	6.33	8.57	3.81		
30	HEMBA1001024	0.31	1.14	2.16	2.87	3.97	1.26		
	HEMBA1001026	0.42	1.52	1.86	2	3.22	2		
	HEMBA1001043	1.43	2.46	2.38	4.63	5.28	4.25	**	+
35	HEMBA1001051	3.36	2.79	11.52	13.26	18.17	18.47	*	+
	HEMBA1001052	0.86	2.15	2.18	1.75	3.58	2.48		
	HEMBA1001059	5.62	9.28	26.25	40.62	56.12	43.49	*	+
	HEMBA1001060	2.66	3.67	6.45	10.78	8.35	9.62	*	+
40	HEMBA1001064	2.12	2.87	3.3	6.04	6.48	4.69	**	+
	HEMBA1001071	29.39	41.54	55.57	143.9	102.43	121.71	**	+
	HEMBA1001077	2.37	1.77	5.21	5.36	6.66	3.96		
	HEMBA1001078	2.18	2.6	5.91	13.3	13.21	11.09	**	+
45	HEMBA1001080	4.03	3.46	11.86	24.15	26.66	26.65	**	+
	HEMBA1001084	1.27	2.37	2.9	5.07	5.88	5.13	**	+
	HEMBA1001085	1.24	2.87	4.04	4.34	5.41	4.56		
50	HEMBA1001088	6.62	6	8.04	3.79	4.34	5.81		
	HEMBA1001093	0.61	1.76	2.72	3.09	3.02	2.99		
	HEMBA1001094	0.64	0.78	2.07	2.08	2.99	1.99		
	HEMBA1001099	1.01	1.72	3	2.5	2.95	2.26		
55	HEMBA1001104	1.2	1.75	2.63	3.64	8.04	3.3		

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	HEMBA1000753	2.56	4.21	6.53	7.98	8.59	4.93		
	HEMBA1000757	1.95	2.95	3.27	6.33	6.68	5.94	**	+
5	HEMBA1000760	3.71	3.81	-6.62	6.96	7.03	6.89		
	HEMBA1000769	1.99	2.36	5.17	3.48	5.87	2.85		
	HEMBA1000773	1	2.32	3.07	2.17	3.18	1.4		
	HEMBA1000774	2.69	2.76	6.37	6.29	7.77	5.22		
10	HEMBA1000780	1.12	2.33	3.66	2.7	4.78	3.29		
	HEMBA1000783	1.32	2.39	4.1	2.78	7.73	2.57		
	HEMBA1000791	2.07	2.4	6.39	4.97	10.17	7.84		
15	HEMBA1000793	12.73	12.73	17.88	19.93	17.49	16.69		
	HEMBA1000802	1.57	1.65	2.59	2.07	4.41	1.1		
	HEMBA1000813	38.24	35.83	34.83	54.63	42.38	53.94	*	+
	HEMBA1000817	2.63	3.82	5.44	5.12	7.02	5.49		
20	HEMBA1000822	1.83	2.89	4.1	4.42	5.76	3.91		
	HEMBA1000827	2.26	2.74	6.45	9.31	7.75	6.94	*	+
	HEMBA1000833	3.1	4.46	7.31	8.06	4.49	4.85		
25	HEMBA1000835	12.53	15.55	75.61	94.51	110.02	86.95	*	+
	HEMBA1000843	1.21	2.2	4.6	3.32	5.63	4.93		
	HEMBA1000851	2.13	1.26	3.5	2.7	5.61	2.74		
	HEMBA1000852	1.95	1.83	5.5	3.52	5.49	3.83		
30	HEMBA1000867	0.85	2.79	4.72	2.77	5.39	3.07		
	HEMBA1000869	0.58	1.29	2.51	2.84	3.97	2.38		
	HEMBA1000870	2.56	2.97	2.59	3.39	5.16	5.49	*	+
35	HEMBA1000872	1.44	2.87	4.01	4.31	4.14	4.34		
	HEMBA1000875	1.89	3.09	5	3.8	4.38	3.77		
	HEMBA1000876	1.75	3.36	4.64	3.9	6.21	4.9		
	HEMBA1000907	1.99	2.47	3.81	3.21	7.15	5.53		
40	HEMBA1000908	0.81	2.06	3.85	2	5.43	1.98		
	HEMBA1000910	1.97	1.61	3.71	3.35	5.25	2.98		
	HEMBA1000918	0.76	1.34	4.37	4.93	6.54	6.95	*	+
	HEMBA1000919	0.86	1.97	2.19	2.49	3.07	3.07		
45	HEMBA1000934	2.5	2.56	1.16	2.14	3.51	2.5		
	HEMBA1000935	1.46	1.62	4.21	2.08	5.15	3.64		
	HEMBA1000940	1.98	3.08	3.1	2.52	9.96	5.72		
50	HEMBA1000942	2.31	2.27	4.77	4.81	7.75	6.69		
	HEMBA1000943	0.58	1.25	2.28	1.83	3.38	2.18		
	HEMBA1000946	3.63	4.04	4.54	6.87	14.9	8.4		
	HEMBA1000960	2.63	3.48	9.97	10.24	12.79	10.7		
55	HEMBA1000962	1.99	2.18	2.01	4.43	3.83	4.56	**	+

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	HEMBA1000575	3.73	4.91	10.84	10.19	15.17	13.08		
	HEMBA1000588	1.75	2.49	4.16	3.12	5.5	3.83		
5	HEMBA1000580	0.59	1.02	2.06	2.24	2.53	1.35		
	HEMBA1000591	3.17	3.3	5.18	10.84	12.16	9.8	**	+
	HEMBA1000592	4.2	5.19	7.77	13.85	14.94	11.78	**	+
10	HEMBA1000594	1.95	1.97	3.16	4	5.86	4.94	*	+
	HEMBA1000604	1.19	3.37	3.48	5.41	10.91	5.29		
	HEMBA1000607	2.83	5.09	12.7	15.52	18.13	20.66	*	+
	HEMBA1000608	0.9	2.34	2.46	2.6	5.5	2.31		
15	HEMBA1000622	0.96	2.19	3.55	3.61	5.24	3.8		
	HEMBA1000634	17.56	22.96	30.36	71.62	60.59	51.59	**	+
	HEMBA1000636	4.59	3.95	6.78	15.48	12.35	12.73	**	+
20	HEMBA1000637	0.93	0.48	2.58	2.42	3.19	2.21		
	HEMBA1000655	1.33	2.11	4.84	6.91	5.57	6.31	*	+
	HEMBA1000657	1.35	1.78	3.24	4.89	5.28	3.26	*	+
	HEMBA1000662	1.3	2.42	2.73	2.52	3.78	2.72		
25	HEMBA1000664	0.94	1.6	2.87	3.11	4.63	2.94		
	HEMBA1000671	2.96	3.84	11.68	21.25	18.69	15.76	*	+
	HEMBA1000673	1.46	2.23	4.76	7.44	7.49	5.51	*	+
	HEMBA1000675	4.18	3.09	4.54	8.18	7.19	8.04	**	+
30	HEMBA1000678	2.23	2.7	4.47	5.03	7.16	5.16		
	HEMBA1000682	3.4	4.64	8.41	13.76	13.69	14.29	**	+
	HEMBA1000686	2.73	3.88	4.83	6.23	6.6	5.32	*	+
35	HEMBA1000702	1.56	2.07	5.25	4.15	5.78	4.32		
	HEMBA1000705	0.65	1.71	3.43	2.34	3.21	1.64		
	HEMBA1000713	3.31	5.6	6.12	6.94	5.86	5.47		
	HEMBA1000718	2.14	2.7	5.25	6.11	5.09	5.95		
40	HEMBA1000719	9.64	12.27	17.77	16.64	15.52	15.64		
	HEMBA1000722	1.97	1.7	3.6	6.55	6.45	5.02	**	+
	HEMBA1000726	2.2	2.23	5.12	9.4	8.77	9.36	**	+
45	HEMBA1000727	4.09	5.35	6.41	5.13	9.08	8.37		
	HEMBA1000732	1.22	2.74	4.21	4.93	5.58	4.42		
	HEMBA1000736	1.56	2.15	3.24	4.11	5.19	4.62	*	+
	HEMBA1000743	1.25	2.72	3.41	5.05	4.88	4.16	*	+
50	HEMBA1000745	1.59	2.47	3.64	4.88	5.33	3.49		
	HEMBA1000747	1.19	1.59	2.56	2.35	3.12	1.49		
	HEMBA1000748	1.67	1.51	4.85	5.11	6.08	4.81		
	HEMBA1000749	1.14	2.04	5.69	5.98	5.91	5.96		
55	HEMBA1000752	1.4	2.3	4.38	3.69	4.53	3.85		

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	HEMBA1000443	1.19	1.9	3.12	2.99	6.28	3.59		
	HEMBA1000446	38.48	43.56	75.05	56.34	60.86	69.87		
5	HEMBA1000456	5.19	4.41	6.5	7.45	5.62	8.77		
	HEMBA1000459	1.95	2.11	4.24	3.46	6.17	5.55		
	HEMBA1000460	7.46	7.84	8.87	13.59	12.54	18.45	*	+
10	HEMBA1000462	2.11	3.51	5.04	6.05	5.16	7.49		
	HEMBA1000464	1.33	0.96	1.73	1.69	2.74	2.53		
	HEMBA1000468	1.25	1.44	2.43	1.69	3.48	2.22		
	HEMBA1000469	2.89	3.37	8.1	5.42	8.81	8.01		
15	HEMBA1000477	2.87	3.03	7.4	5.41	9.68	6.83		
	HEMBA1000481	29.67	31.97	31.95	42.76	52.75	25.82		
	HEMBA1000488	1.75	2.43	2.96	3.11	5.9	3		
20	HEMBA1000490	1.34	2	3.49	4.41	3.7	2.88		
	HEMBA1000491	1.21	1.71	2.85	4.24	4.99	5.97	*	+
	HEMBA1000498	2.12	3.21	4.55	4.39	7.76	5.94		
	HEMBA1000501	2.22	3.36	6.25	6.44	8.93	9.74	*	+
25	HEMBA1000504	2.93	3.18	4.82	3.63	5.37	3.83		
	HEMBA1000505	0.81	1.97	3.33	2.72	5.1	3.58		
	HEMBA1000507	1.02	2.24	5.29	4.17	8.62	7		
	HEMBA1000508	2.25	2.3	7.65	4.84	8.57	6.64		
30	HEMBA1000518	1.38	0.96	0.98	1.89	2.97	1.8	*	+
	HEMBA1000519	9.5	7.28	15.97	19.28	20.99	19.72	*	+
	HEMBA1000520	0.45	1.12	1.18	1.94	4.83	4.3	*	+
35	HEMBA1000523	2.32	1.88	3.22	3.48	5.33	3.65		
	HEMBA1000531	1.39	1.46	2.44	2.67	5.34	4.63	*	+
	HEMBA1000534	0.55	0.95	2.97	6.63	11.62	10.39	**	+
	HEMBA1000538	0.51	1.08	2.31	12.58	21.02	13.18	**	+
40	HEMBA1000540	2.8	3.11	6.06	5.82	10.38	6.39		
	HEMBA1000542	9.16	7.79	43.94	62.25	95.7	81.15	*	+
	HEMBA1000545	1.51	2.31	1.65	3.19	4.29	3.7	**	+
	HEMBA1000547	2.99	3.12	4.94	4.94	5.3	4.97		
45	HEMBA1000551	2.32	1.99	9.54	4.68	7.33	9.81		
	HEMBA1000555	3.81	3.23	6.39	5.03	6.43	8.08		
	HEMBA1000557	2.16	2.06	6.07	3.98	6.46	5.06		
50	HEMBA1000561	1.71	2.9	4.9	1.63	4.39	3.67		
	HEMBA1000563	1.73	1.85	4.09	2.72	3.94	2.83		
	HEMBA1000567	1.02	1.01	1.67	1.21	2.59	1.92		
	HEMBA1000568	2.19	2.5	6.09	7.62	6.65	6.84		
55	HEMBA1000569	1.3	2.8	3.02	2.18	6.47	2.3		

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	HEMBA1000269	1.9	2.34	3.69	4.41	4.09	2.51		
	HEMBA1000275	5.31	4.29	8.03	7.96	12.04	8.54		
5	HEMBA1000280	1.43	0.83	2.19	3.3	4.08	4	**	+
	HEMBA1000282	1.15	1.01	4.23	6.29	7.01	5.46	*	+
	HEMBA1000287	2.86	3.19	4.45	5.81	6.04	6.37	**	+
	HEMBA1000288	1.37	2.23	6.13	3.51	6.02	3.85		
10	HEMBA1000290	1.01	2.17	4.11	2.46	3.26	2.73		
	HEMBA1000296	2.4	3.66	5.49	6.15	6.55	5.84		
	HEMBA1000300	1.22	2.73	6.6	7.64	8.88	7.23		
15	HEMBA1000302	0.93	2.17	2.86	3.04	3.74	1.97		
	HEMBA1000303	1.36	2.15	3.57	4.13	4.43	3		
	HEMBA1000304	1.06	1.99	4.26	5.51	7.28	4.87	*	+
	HEMBA1000307	1.21	1.73	2.65	4.4	5.64	2.99	*	+
20	HEMBA1000312	6	8.7	10.77	13.2	9.18	9.65		
	HEMBA1000318	1.5	4.22	3.25	5.39	6.05	4.49		
	HEMBA1000327	2.18	3.7	3.34	10.58	6.06	6.02	*	+
25	HEMBA1000333	0.68	2.75	4.33	3.12	4.74	2.98		
	HEMBA1000338	1.61	2.84	5.33	5.8	5.78	4.32		
	HEMBA1000343	1.79	3.5	3.69	5.55	6.7	3.99		
	HEMBA1000349	0.97	1.52	3.24	3.9	5.37	4.09	*	+
30	HEMBA1000351	1.6	2.06	5.75	4.8	6.22	5.24		
	HEMBA1000355	1.52	3.09	4.09	3.78	5.14	3.59		
	HEMBA1000356	9.3	10.42	14.39	26.93	22.26	24.97	**	+
35	HEMBA1000357	1.88	2.11	4.76	3.81	5.7	4.62		
	HEMBA1000366	1.67	1.94	3.83	3.14	4.75	3.28		
	HEMBA1000369	1.87	2.94	5.17	2.82	5.2	4.56		
	HEMBA1000370	2.45	3.4	4.63	3.75	5.34	3.6		
40	HEMBA1000376	3.64	4.55	14.48	26.69	29.98	28.36	**	+
	HEMBA1000387	2.95	3.19	6.2	7.85	7.62	8.15	*	+
	HEMBA1000389	2.88	3.74	8.83	14.4	10.9	13.61	*	+
45	HEMBA1000390	1.86	2.27	3.5	4.28	4.98	3.95	*	+
	HEMBA1000392	1.49	1.4	3.06	2.58	3.78	1.94		
	HEMBA1000396	1.82	2.16	3.45	3.43	4.93	3.34		
	HEMBA1000411	1.01	1.41	4.49	1.94	4.41	2.21		
50	HEMBA1000418	2.85	3.21	4.41	7.75	6.81	5.17	*	+
	HEMBA1000422	0.99	1.89	2.14	2.64	4.03	2.89		
	HEMBA1000428	0.36	2.43	3.09	2.58	3.31	2.75		
	HEMBA1000434	0.54	2.19	2.93	2.11	3.6	2.69		
55	HEMBA1000442	0.82	2.2	3.37	2.13	3.8	2.28		

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	HEMBA1000047	1.17	1.99	3.83	2.98	4.47	3.78		
	HEMBA1000048	3.76	4.75	3.64	8.73	12.38	5.48		
5	HEMBA1000050	0.48	1.77	1.78	1.4	3.39	1.46		
	HEMBA1000053	1.64	1.28	2.57	4.68	4.03	3.99	**	+
	HEMBA1000060	1.88	2.71	4.51	7.29	7.94	9.74	**	+
10	HEMBA1000072	52.79	53.46	135.73	165.97	221.75	230.97	*	+
	HEMBA1000073	16.54	11.43	27.32	22.4	36.09	33.78		
	HEMBA1000076	5.06	5.33	9.77	12.16	10.46	11.15	*	+
	HEMBA1000084	4.75	4.46	20.71	30.15	43.67	33.92	*	+
15	HEMBA1000087	0.51	1	3.32	0.65	2.82	1.61		
	HEMBA1000088	1.98	2.97	4.6	6.2	9.87	8.46	*	+
	HEMBA1000091	6.36	5.4	17.56	30.15	44.04	35.43	**	+
	HEMBA1000111	1.52	1.77	3.63	5.29	6.65	6.4	**	+
20	HEMBA1000121	0.86	1.17	3.58	3.52	4.47	5.83		
	HEMBA1000128	1.52	2.99	6.04	4.28	6.05	5.93		
	HEMBA1000129	2.04	1.81	3.95	2.66	3.26	3.32		
25	HEMBA1000141	2.31	3.45	5.98	3.56	6.67	5.6		
	HEMBA1000146	0.84	1.29	2.96	1.93	4.98	3.52		
	HEMBA1000150	3.34	3.29	10.65	8.27	11.97	9.46		
	HEMBA1000154	25.17	29.21	82.33	128.3	134.42	139.59	**	+
30	HEMBA1000156	3.28	4	5.87	8.69	6.19	6.13		
	HEMBA1000158	7.98	10.04	12.52	16.99	15.47	12.88		
	HEMBA1000168	1.21	2.2	4.11	5.7	7.3	5.21	*	+
35	HEMBA1000180	0.4	2.04	2.87	2.86	4.06	2.05		
	HEMBA1000185	1.65	3.84	4.88	7.5	9.46	9.07	**	+
	HEMBA1000188	1.37	1.64	3.31	4.94	4.19	3.35		
	HEMBA1000193	1.53	0.66	3.16	2.68	4.33	2.5		
40	HEMBA1000194	2.18	2.95	5.68	9.11	8.74	8.83	**	+
	HEMBA1000201	2.6	4.47	9.74	13.45	14.8	14.65	*	+
	HEMBA1000213	1.33	1.95	2.76	2.08	4.49	3.7		
45	HEMBA1000216	1.26	1.82	3.27	2.92	5.2	3.47		
	HEMBA1000227	0.99	2.27	2.38	3.28	4.21	1.83		
	HEMBA1000231	1.5	1.97	5	7.56	7.19	6.16	*	+
	HEMBA1000237	4.5	6.13	9.14	14.79	18.3	14.71	**	+
50	HEMBA1000243	0.6	1.89	3.01	4.3	4.34	3.67	*	+
	HEMBA1000244	1.54	2.45	3.78	6.08	5.58	3.36		
	HEMBA1000251	1.15	1.92	2.97	2.26	4.59	3.22		
	HEMBA1000254	0.69	1.8	4.81	3.57	4.97	3.58		
55	HEMBA1000264	0.84	2.28	3.01	2.84	3.23	3.12		

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	BNGH41000198	1.32	2.3	4.35	2.06	2.55	2.22		
	BNGH41000219	2.29	3.91	5.61	12.4	13.73	10.76	**	+
5	BNGH41000229	9.65	9.99	12.99	18.34	18.92	18.94	**	+
	BNGH41000237	8.4	12.99	12.61	27.63	11.26	13.45		
	BNGH41000238	1.56	2.59	6.77	3.45	4.55	3.32		
10	BNGH41000243	5.56	8.95	6.71	15.03	12.55	16.36	**	+
	BNGH41000270	2.94	2.77	2.88	3.67	3.99	3.74	**	+
	BRAWH1000004	1	2.19	6.99	6.45	8.36	6		
	BRAWH1000018	1.8	2.24	5.06	4.43	6.95	5.24		
15	BRAWH1000021	1.33	2.73	4.81	4.16	5.85	5.21		
	BRAWH1000027	0.58	1.7	1.62	2.39	3.65	2.63	*	+
	BRAWH1000029	2.32	3.63	6.21	6.03	6.73	4.81		
20	BRAWH1000040	4.68	4.98	8.01	7.28	7.2	8.67		
	BRAWH1000050	11.04	10.47	43.79	51.7	73.7	60.92	*	+
	BRAWH1000051	2.14	0.63	2.71	2.25	4.43	1.04		
	BRAWH1000060	7.84	8.07	48.26	59.16	66.12	63.86	*	+
25	BRAWH1000075	1.85	1.86	2.98	2.07	4.4	2.34		
	BRAWH1000081	1.88	2.78	7.19	5.9	10.82	7.4		
	BRAWH1000084	30.23	30.57	65.21	235.81	180.86	211.35	**	+
30	BRAWH1000095	1.38	2.47	4.51	3	4.78	2.67		
	BRAWH1000096	1.37	2.89	4.71	3.7	4.8	5.17		
	BRAWH1000097	3.32	3.27	10.74	9.24	10.62	7.75		
	BRAWH1000100	4.77	5.19	7.69	6.98	7.06	7.28		
35	BRAWH1000101	12	12.04	36.52	46.19	41.09	50.21	*	+
	BRAWH1000104	1.37	0.92	4.33	1.47	4.47	2.41		
	BRAWH1000107	0.62	1.88	2.48	2.43	5.03	3.15		
	BRAWH1000110	4.4	4.06	16.81	13.87	11.1	15.74		
40	BRAWH1000111	3.98	6.14	6.05	8.85	8.95	10.64	*	+
	BRAWH1000135	4.95	4.91	7.7	7.37	9.42	9.98		
	BRAWH1000190	2.22	3.84	5.07	4.66	7.16	4.99		
45	HEMBA1000005	5.91	6.44	11.97	17.55	22.88	18.65	*	+
	HEMBA1000006	2.61	3.17	4.64	3.08	8.49	4.75		
	HEMBA1000012	10.97	11.75	51.07	71.4	106.82	74.8	*	+
	HEMBA1000020	50.65	49.12	113.3	197.41	293.79	216.89	*	+
50	HEMBA1000030	1.93	3.08	4.67	5.72	3.62	6.43		
	HEMBA1000034	3.27	3.21	5.35	4.62	10.29	6.85		
	HEMBA1000042	1.64	3.17	6	4.72	6.92	8.12		
	HEMBA1000045	7.13	9.44	11.07	9.55	14.43	10.44		
55	HEMBA1000046	1.14	2.24	2.77	3.73	5.3	4.34	*	+

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	GAPDH (Cr1)	0.4	0.8	0.89	0.9	1	1.15		
	β actin (Cr2)	385.94	262.23	582.98	443.28	422.61	573.47		
5	ADRGL1000005	2.72	2.97	4.46	7.27	7.45	3.51		
	ADRGL1000007	4.36	5.19	9.58	20.78	19.59	18.29	**	+
	ADRGL1000009	0.99	1.25	1.64	2.16	4.08	2.02		
10	ADRGL1000011	1.98	3.56	5.24	22.22	23.49	19.81	**	+
	ADRGL1000027	0.79	1.22	1.66	2.82	4.99	1.9		
	ADRGL1000058	4.12	7.08	26.9	62.55	67.32	49.15	**	+
	ADRGL1000069	1.91	1.68	2.47	14.19	14.54	13.74	**	+
15	ADRGL1000077	1.98	2	2.54	5.5	2.9	4.16		
	ADRGL1000092	2.99	4.79	12.53	21.46	22.09	26.19	**	+
	ADRGL1000099	2.77	4.79	12.85	23.61	24.02	25.56	**	+
20	ADRGL1000136	20.49	27.18	31.85	62.44	40.69	48.29	*	+
	ADRGL1000147	2.09	2.58	5.47	5.69	7.52	3.85		
	ADRGL1000159	1.51	1.77	3.07	3.4	4.71	2.59		
	ADRGL1000160	2.42	4.34	6.89	8.08	7.24	7.06		
25	ADRGL1000171	0.95	1.11	1.64	1.89	2.69	1.87		
	ADRGL1000181	0.64	1.37	1.74	3.99	4.27	3.89	**	+
	BGG111000015	2.13	3.89	5.02	10.49	11.35	9.14	**	+
30	BGG111000016	27.77	35.71	52.17	57.18	48.51	63.57		
	BGG111000017	1.29	3.19	3.14	3.24	3.65	2.34		
	BGG111000022	4.72	4.45	6.75	10.71	5.56	8.27		
	BGG111000031	4.47	6.58	8.77	14.79	11.63	10.04	*	+
35	BGG111000042	9.55	11.29	20.54	23.39	18.75	20.23		
	BGG111000046	8.56	9.77	17.04	34.24	30.76	25.79	**	+
	BNGH41000020	246.16	211.77	380.83	658.32	647.37	559.16	**	+
40	BNGH41000025	4.31	3.12	6.92	11.4	13.1	15.01	**	+
	BNGH41000026	2.71	4.77	7.53	4.45	7.17	6.23		
	BNGH41000027	11.52	13.5	12.69	20.62	12.48	24.91		
	BNGH41000035	23.02	25.91	36.46	51.05	31.83	41.67		
45	BNGH41000037	2.7	5.21	6.72	12.95	8.98	8.59	*	+
	BNGH41000042	14.55	16.06	22.84	49.62	37.57	36.25	**	+
	BNGH41000048	3.92	6.27	25.68	66.19	74.4	66.21	**	+
	BNGH41000056	0.74	1.75	3.26	5.28	7.34	3.75	*	+
50	BNGH41000087	3.36	4.08	5.19	5.59	8.15	3.01		
	BNGH41000091	0.18	1.45	2.47	2.72	3.4	2.14		
	BNGH41000157	6.93	7.99	6.23	13.37	10.28	9.98	*	+
55	BNGH41000169	1.09	1.53	2.99	2.77	4.23	2.59		
	BNGH41000181	3.5	4.06	7.5	5.71	6.81	6.09		

Table 352

Expression of each cDNA in synovial cells or in the synovial cells in the presence of TNF
(This table also contains clones without description in Examples)

In the table, Synoviocyte and Synoviocyte_TNF represent synovial cells and TNF-treated synovial cells, respectively. The assay was performed in triplicate (n=3), and each result is shown in the column of exp.1, exp.2, or exp.3. In addition, "t-test vs TNF" represents a result of test for significance of difference between the untreated synovial cells and the TNF-treated synovial cells. The increase and decrease in the expression level of a particular gene in response to TNF are represented by + and -, respectively. The results of test for significance of difference are shown in the columns of *:p<0.05 and **:p<0.01.

Clone	Synoviocyte			Synoviocyte_TNF			t test INC.	
	exp. 1	exp. 2	exp. 3	exp. 1	exp. 2	exp. 3	vs TNF	and DEC.

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	NT2RP2003339	C-NT2RP2003339	18992	
	NT2RP3001282	C-NT2RP3001282	18993	18994
5	PLACE1001761	C-PLACE1001761.....	18995	18996
	PLACE1004491	C-PLACE1004491	18997	
	PLACE1004686	C-PLACE1004686	18998	
10	PLACE1005574	C-PLACE1005574	18999	
	PLACE1006382	C-PLACE1006382	19000	
	PLACE1006792	C-PLACE1006792	19001	
15	PLACE3000455	C-PLACE3000455	19002	19003
	PLACE4000230	C-PLACE4000230	19004	19005
	THYR01000916	C-THYR01000916	19006	
20	HEMBA1000327	C-HEMBA1000327	19007	19008
	HEMBB1000637	C-HEMBB1000637	19009	19010
	HEMBB1001967	C-HEMBB1001967	19011	
	MAMMA1000266	C-MAMMA1000266	19012	
25	NT2RP2002979	C-NT2RP2002979	19013	
	PLACE1007866	C-PLACE1007866	19014	19015
	PLACE3000350	C-PLACE3000350	19016	19017
30	PLACE4000156	C-PLACE4000156	19018	19019
	THYR01001637	C-THYR01001637	19020	19021
	MAMMA1002215	C-MAMMA1002215	19022	19023
35	MAMMA1002721	C-MAMMA1002721	19024	
	NT2RP2002070	C-NT2RP2002070	19025	

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	THYR01001828	C-THYR01001828	18936	18937
	Y79AA1000346	C-Y79AA1000346	18938	18939
5	Y79AA1001167	C-Y79AA1001167	18940	
	Y79AA1001384	C-Y79AA1001384	18941	18942
	Y79AA1001875	C-Y79AA1001875	18943	
	Y79AA1002103	C-Y79AA1002103	18944	18945
10	HEMBA1000290	C-HEMBA1000290	18946	
	HEMBA1001196	C-HEMBA1001196	18947	18948
	HEMBA1006650	C-HEMBA1006650	18949	18950
	HEMBA1006796	C-HEMBA1006796	18951	18952
15	HEMBB1000337	C-HEMBB1000337	18953	18954
	HEMBB1001619	C-HEMBB1001619	18955	
	MAMMA1000270	C-MAMMA1000270	18956	
20	MAMMA1000559	C-MAMMA1000559	18957	
	MAMMA1000940	C-MAMMA1000940	18958	
	MAMMA1002545	C-MAMMA1002545	18959	
	MAMMA1002972	C-MAMMA1002972	18960	18961
25	NT2RP2001440	C-NT2RP2001440	18962	18963
	NT2RP3002770	C-NT2RP3002770	18964	18965
	NT2RP3003138	C-NT2RP3003138	18966	18967
30	NT2RP3004470	C-NT2RP3004470	18968	
	OVARC1000891	C-OVARC1000891	18969	
	PLACE1001545	C-PLACE1001545	18970	18971
	PLACE1003383	C-PLACE1003383	18972	
35	PLACE1005549	C-PLACE1005549	18973	18974
	PLACE1008455	C-PLACE1008455	18975	
	PLACE4000131	C-PLACE4000131	18976	18977
	PLACE4000261	C-PLACE4000261	18978	18979
40	THYR01001602	C-THYR01001602	18980	
	HEMBA1006092	C-HEMBA1006092	18981	
	HEMBA1006406	C-HEMBA1006406	18982	
45	HEMBB1000790	C-HEMBB1000790	18983	
	HEMBB1000917	C-HEMBB1000917	18984	
	HEMBB1002280	C-HEMBB1002280	18985	
	MAMMA1000802	C-MAMMA1000802	18986	
50	MAMMA1001322	C-MAMMA1001322	18987	
	MAMMA1002597	C-MAMMA1002597	18988	
	MAMMA1002868	C-MAMMA1002868	18989	
55	NT2RP2003161	C-NT2RP2003161	18990	18991

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	PLACE3000124	C-PLACE3000124	18869	18870
	PLACE3000158	C-PLACE3000158	18871	
5	PLACE3000207	C-PLACE3000207	18872	
	PLACE3000221	C-PLACE3000221	18873	18874
	PLACE3000271	C-PLACE3000271	18875	18876
	PLACE3000304	C-PLACE3000304	18877	
10	PLACE3000322	C-PLACE3000322	18878	18879
	PLACE3000341	C-PLACE3000341	18880	
	PLACE3000373	C-PLACE3000373	18881	18882
15	PLACE3000399	C-PLACE3000399	18883	18884
	PLACE3000401	C-PLACE3000401	18885	
	PLACE3000402	C-PLACE3000402	18886	
	PLACE3000406	C-PLACE3000406	18887	18888
20	PLACE3000475	C-PLACE3000475	18889	
	PLACE4000063	C-PLACE4000063	18890	18891
	PLACE4000093	C-PLACE4000093	18892	
	PLACE4000100	C-PLACE4000100	18893	18894
25	PLACE4000247	C-PLACE4000247	18895	18896
	PLACE4000250	C-PLACE4000250	18897	18898
	PLACE4000252	C-PLACE4000252	18899	18900
30	PLACE4000259	C-PLACE4000259	18901	18902
	PLACE4000320	C-PLACE4000320	18903	
	PLACE4000344	C-PLACE4000344	18904	
	PLACE4000367	C-PLACE4000367	18905	18906
35	PLACE4000401	C-PLACE4000401	18907	18908
	PLACE4000411	C-PLACE4000411	18909	18910
	PLACE4000487	C-PLACE4000487	18911	18912
	PLACE4000494	C-PLACE4000494	18913	18914
40	PLACE4000521	C-PLACE4000521	18915	18916
	PLACE4000548	C-PLACE4000548	18917	18918
	SKNMC1000013	C-SKNMC1000013	18919	18920
45	SKNMC1000091	C-SKNMC1000091	18921	18922
	THYRO1000343	C-THYRO1000343	18923	18924
	THYRO1000569	C-THYRO1000569	18925	18926
	THYRO1001142	C-THYRO1001142	18927	
50	THYRO1001189	C-THYRO1001189	18928	18929
	THYRO1001320	C-THYRO1001320	18930	18931
	THYRO1001537	C-THYRO1001537	18932	18933
55	THYRO1001721	C-THYRO1001721	18934	18935

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	PLACE1001608	C-PLACE1001608	18803	18804
	PLACE1002004	C-PLACE1002004	18805	18806
5	PLACE1002256	C-PLACE1002256	18807	
	PLACE1002437	C-PLACE1002437	18808	18809
	PLACE1002591	C-PLACE1002591	18810	18811
	PLACE1002665	C-PLACE1002665	18812	18813
10	PLACE1003864	C-PLACE1003864	18814	18815
	PLACE1004793	C-PLACE1004793	18816	18817
	PLACE1004913	C-PLACE1004913	18818	18819
	PLACE1004979	C-PLACE1004979	18820	
15	PLACE1005052	C-PLACE1005052	18821	18822
	PLACE1005055	C-PLACE1005055	18823	18824
	PLACE1005128	C-PLACE1005128	18825	18826
20	PLACE1005162	C-PLACE1005162	18827	18828
	PLACE1005176	C-PLACE1005176	18829	18830
	PLACE1005467	C-PLACE1005467	18831	18832
	PLACE1005584	C-PLACE1005584	18833	18834
25	PLACE1005611	C-PLACE1005611	18835	18836
	PLACE1005802	C-PLACE1005802	18837	
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30	PLACE1005898	C-PLACE1005898	18839	18840
	PLACE1005932	C-PLACE1005932	18841	
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35	PLACE1006795	C-PLACE1006795	18845	
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	PLACE1007557	C-PLACE1007557	18848	
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40	PLACE1008181	C-PLACE1008181	18851	
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45	PLACE1009935	C-PLACE1009935	18856	18857
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	PLACE2000170	C-PLACE2000170	18866	
55	PLACE2000335	C-PLACE2000335	18867	18868

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	NT2RP3004125	C-NT2RP3004125	18744	18745
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20	NT2RP4000515	C-NT2RP4000515	18760	
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	NT2RP4000519	C-NT2RP4000519	18762	18763
25	NT2RP5003512	C-NT2RP5003512	18764	18765
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	OVARC1000802	C-OVARC1000802	18771	
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35	OVARC1001072	C-OVARC1001072	18776	18777
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55	PLACE1001366	C-PLACE1001366	18801	18802

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	NT2RP3002818	C-NT2RP3002818	18687	18688
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	NT2RP3003059	C-NT2RP3003059	18692	18693
	NT2RP3003121	C-NT2RP3003121	18694	18695
	NT2RP3003133	C-NT2RP3003133	18696	18697
25	NT2RP3003155	C-NT2RP3003155	18698	18699
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30	NT2RP3003264	C-NT2RP3003264	18704	18705
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	NT2RM1000702	C-NT2RM1000702	18603	18604
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	NT2RM1001082	C-NT2RM1001082	18609	
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	NT2RM2001360	C-NT2RM2001360	18614	18615
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	NT2RM2001803	C-NT2RM2001803	18618	18619
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	NT2RP2002105	C-NT2RP2002105	18636	18637
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	NT2RP2002843	C-NT2RP2002843	18642	
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	NT2RP2003799	C-NT2RP2003799	18647	
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	MAMMA1002617	C-MAMMA1002617	18557	18558
	MAMMA1002618	C-MAMMA1002618	18559	
	MAMMA1002636	C-MAMMA1002636	18560	
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	MAMMA1002665	C-MAMMA1002665	18563	
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	MAMMA1002880	C-MAMMA1002880	18575	18576
35	MAMMA1002892	C-MAMMA1002892	18577	
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	MAMMA1003031	C-MAMMA1003031	18588	
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	NT2RM1000341	C-NT2RM1000341	18595	18596
	NT2RM1000539	C-NT2RM1000539	18597	18598
55	NT2RM1000553	C-NT2RM1000553	18599	18600

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	HEMBB1002534	C-HEMBB1002534	18495	
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45	MAMMA1001630	C-MAMMA1001630	18530	
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35	HEMBA1000835	C-HEMBA1000835	18451	18452
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40	HEMBA1001500	C-HEMBA1001500	18458	18459
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	HEMBA1001872	C-HEMBA1001872	18474	18475
55	HEMBA1001905	C-HEMBA1001905	18476	18477

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5	Y79AA1001866	C-Y79AA1001866	18347	18348
	Y79AA1001874	C-Y79AA1001874	18349	18350
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	Y79AA1002220	C-Y79AA1002220	18357	
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	Y79AA1002258	C-Y79AA1002258	18360	18361
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	Y79AA1002416	C-Y79AA1002416	18366	18367
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40	HEMBA1003838	C-HEMBA1003838	18394	
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45	HEMBA1004086	C-HEMBA1004086	18398	
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	HEMBA1004353	C-HEMBA1004353	18401	
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50	HEMBA1004499	C-HEMBA1004499	18404	18405
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	HEMBA1004638	C-HEMBA1004638	18408	18409
55	HEMBA1004669	C-HEMBA1004669	18410	18411

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	THYR01001347	C-THYR01001347	18286	18287
	THYR01001363	C-THYR01001363	18288	
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15	THYR01001403	C-THYR01001403	18291	18292
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	THYR01001487	C-THYR01001487	18302	
25	THYR01001584	C-THYR01001584	18303	18304
	THYR01001661	C-THYR01001661	18305	18306
	THYR01001746	C-THYR01001746	18307	
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30	THYR01001854	C-THYR01001854	18309	
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	Y79AA1000033	C-Y79AA1000033	18312	18313
	Y79AA1000231	C-Y79AA1000231	18314	18315
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45	Y79AA1000966	C-Y79AA1000966	18329	18330
	Y79AA1000969	C-Y79AA1000969	18331	
	Y79AA1000985	C-Y79AA1000985	18332	18333
	Y79AA1001061	C-Y79AA1001061	18334	
50	Y79AA1001068	C-Y79AA1001068	18335	18336
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	Y79AA1001299	C-Y79AA1001299	18339	18340
55	Y79AA1001511	C-Y79AA1001511	18341	18342

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	SKNMC1000050	C-SKNMC1000050	18209	
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	THYR01000092	C-THYR01000092	18216	18217
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	THYR01000124	C-THYR01000124	18220	18221
	THYR01000197	C-THYR01000197	18222	18223
15	THYR01000199	C-THYR01000199	18224	18225
	THYR01000206	C-THYR01000206	18226	
	THYR01000242	C-THYR01000242	18227	18228
	THYR01000253	C-THYR01000253	18229	18230
20	THYR01000270	C-THYR01000270	18231	18232
	THYR01000288	C-THYR01000288	18233	18234
	THYR01000320	C-THYR01000320	18235	
	THYR01000358	C-THYR01000358	18236	18237
25	THYR01000368	C-THYR01000368	18238	18239
	THYR01000381	C-THYR01000381	18240	18241
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	THYR01000988	C-THYR01000988	18270	18271
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20	NT2RP3003203	C-NT2RP3003203	17591	17592
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25	NT2RP3003242	C-NT2RP3003242	17598	17599
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30	NT2RP3003409	C-NT2RP3003409	17606	17607
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	NT2RP3003543	C-NT2RP3003543	17610	17611
35	NT2RP3003552	C-NT2RP3003552	17612	17613
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40	NT2RP3003621	C-NT2RP3003621	17620	17621
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	NT2RP3003656	C-NT2RP3003656	17624	17625
45	NT2RP3003659	C-NT2RP3003659	17626	17627
	NT2RP3003686	C-NT2RP3003686	17628	17629
	NT2RP3003701	C-NT2RP3003701	17630	17631
	NT2RP3003716	C-NT2RP3003716	17632	17633
50	NT2RP3003726	C-NT2RP3003726	17634	17635
	NT2RP3003795	C-NT2RP3003795	17636	17637
	NT2RP3003805	C-NT2RP3003805	17638	17639
55	NT2RP3003809	C-NT2RP3003809	17640	17641

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5	NT2RP3001554	C-NT2RP3001554.....	17498	17499
	NT2RP3001712	C-NT2RP3001712	17500	17501
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10	NT2RP3001730	C-NT2RP3001730	17506	17507
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15	NT2RP3001857	C-NT2RP3001857	17512	17513
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	NT2RP3002351	C-NT2RP3002351	17537	17538
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	NT2RP3002545	C-NT2RP3002545	17543	17544
40	NT2RP3002549	C-NT2RP3002549	17545	17546
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	NT2RP3002603	C-NT2RP3002603	17549	17550
	NT2RP3002628	C-NT2RP3002628	17551	17552
45	NT2RP3002659	C-NT2RP3002659	17553	17554
	NT2RP3002660	C-NT2RP3002660	17555	17556
	NT2RP3002682	C-NT2RP3002682	17557	17558
	NT2RP3002687	C-NT2RP3002687	17559	17560
50	NT2RP3002688	C-NT2RP3002688	17561	17562
	NT2RP3002701	C-NT2RP3002701	17563	17564
	NT2RP3002785	C-NT2RP3002785	17565	17566
55	NT2RP3002869	C-NT2RP3002869	17567	17568

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	NT2RP2005496	C-NT2RP2005496	17432	
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	NT2RP2005645	C-NT2RP2005645	17437	
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	NT2RP2005741	C-NT2RP2005741	17442	
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	NT2RP2005882	C-NT2RP2005882	17449	
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	NT2RP2006103	C-NT2RP2006103	17452	17453
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	NT2RP2006258	C-NT2RP2006258	17456	
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	NT2RP3000487	C-NT2RP3000487	17470	17471
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	NT2RP3001245	C-NT2RP3001245	17489	
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55	NT2RP3001356	C-NT2RP3001356	17492	17493

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	NT2RP2004709	C-NT2RP2004709	17387	
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25	NT2RP2004767	C-NT2RP2004767	17392	
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30	NT2RP2004962	C-NT2RP2004962	17397	
	NT2RP2004982	C-NT2RP2004982	17398	
	NT2RP2005003	C-NT2RP2005003	17399	17400
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	NT2RP2005031	C-NT2RP2005031	17406	17407
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	NT2RP2005168	C-NT2RP2005168	17412	17413
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45	NT2RP2005325	C-NT2RP2005325	17416	17417
	NT2RP2005336	C-NT2RP2005336	17418	17419
	NT2RP2005344	C-NT2RP2005344	17420	17421
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	NT2RP2005457	C-NT2RP2005457	17427	17428
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	NT2RP2003293	C-NT2RP2003293	17309	
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	NT2RP2003401	C-NT2RP2003401	17312	17313
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	NT2RP2003581	C-NT2RP2003581	17322	17323
	NT2RP2003643	C-NT2RP2003643	17324	17325
25	NT2RP2003702	C-NT2RP2003702	17326	17327
	NT2RP2003704	C-NT2RP2003704	17328	17329
	NT2RP2003727	C-NT2RP2003727	17330	17331
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	NT2RP2003781	C-NT2RP2003781	17334	17335
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	NT2RP2003871	C-NT2RP2003871	17338	17339
35	NT2RP2003885	C-NT2RP2003885	17340	
	NT2RP2003912	C-NT2RP2003912	17341	17342
	NT2RP2003976	C-NT2RP2003976	17343	
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	NT2RP2004098	C-NT2RP2004098	17347	17348
	NT2RP2004142	C-NT2RP2004142	17349	17350
45	NT2RP2004170	C-NT2RP2004170	17351	17352
	NT2RP2004194	C-NT2RP2004194	17353	17354
	NT2RP2004207	C-NT2RP2004207	17355	
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	NT2RP2004242	C-NT2RP2004242	17360	17361
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55	NT2RP2004321	C-NT2RP2004321	17364	

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	NT2RP2001839	C-NT2RP2001839	17238	17239
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	NT2RP2001946	C-NT2RP2001946	17251	17252
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	NT2RP2002033	C-NT2RP2002033	17255	
	NT2RP2002041	C-NT2RP2002041	17256	17257
	NT2RP2002047	C-NT2RP2002047	17258	
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	NT2RP2002741	C-NT2RP2002741	17287	17288
	NT2RP2002752	C-NT2RP2002752	17289	
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	NT2RP2002857	C-NT2RP2002857	17292	17293
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55	NT2RP2003073	C-NT2RP2003073	17296	

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	NT2RM4002571	C-NT2RM4002571	17164	17165
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	NT2RP1000522	C-NT2RP1000522	17168	17169
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	NT2RP1000677	C-NT2RP1000677	17172	17173
	NT2RP1000701	C-NT2RP1000701	17174	17175
	NT2RP1000834	C-NT2RP1000834	17176	17177
15	NT2RP1000860	C-NT2RP1000860	17178	17179
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25	NT2RP1001177	C-NT2RP1001177	17190	17191
	NT2RP1001185	C-NT2RP1001185	17192	
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	NT2RP2000198	C-NT2RP2000198	17204	17205
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	NT2RP2001634	C-NT2RP2001634	17224	17225
55	NT2RP2001660	C-NT2RP2001660	17226	17227

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	NT2RM2001637	C-NT2RM2001637	17100	17101
	NT2RM2001641	C-NT2RM2001641	17102	
15	NT2RM2001670	C-NT2RM2001670	17103	17104
	NT2RM2001699	C-NT2RM2001699	17105	17106
	NT2RM2001706	C-NT2RM2001706	17107	
	NT2RM2001718	C-NT2RM2001718	17108	17109
20	NT2RM2001727	C-NT2RM2001727	17110	17111
	NT2RM2001805	C-NT2RM2001805	17112	17113
	NT2RM4000086	C-NT2RM4000086	17114	17115
25	NT2RM4000215	C-NT2RM4000215	17116	17117
	NT2RM4000414	C-NT2RM4000414	17118	
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30	NT2RM4000657	C-NT2RM4000657	17123	17124
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35	NT2RM4000996	C-NT2RM4000996	17130	17131
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40	NT2RM4001569	C-NT2RM4001569	17136	17137
	NT2RM4001819	C-NT2RM4001819	17138	
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	NT2RM4001938	C-NT2RM4001938	17141	17142
45	NT2RM4002062	C-NT2RM4002062	17143	17144
	NT2RM4002073	C-NT2RM4002073	17145	17146
	NT2RM4002093	C-NT2RM4002093	17147	17148
	NT2RM4002109	C-NT2RM4002109	17149	17150
50	NT2RM4002146	C-NT2RM4002146	17151	17152
	NT2RM4002194	C-NT2RM4002194	17153	17154
	NT2RM4002390	C-NT2RM4002390	17155	17156
55	NT2RM4002398	C-NT2RM4002398	17157	17158

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	MAMMA1002322	C-MAMMA1002322	17034	
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35	MAMMA1002699	C-MAMMA1002699	17063	17064
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50	MAMMA1003047	C-MAMMA1003047	17078	17079
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	NT2RM1000018	C-NT2RM1000018	17082	17083
55	NT2RM1000037	C-NT2RM1000037	17084	17085

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35	HEMBA1006976	C-HEMBA1006976	16601	16602
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50	HEMBA1007151	C-HEMBA1007151	16615	16616
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	HEMBA1006173	C-HEMBA1006173	16555	16556
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	Y79AA1002208	C-Y79AA1002208	16089	16090
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	Y79AA1002246	C-Y79AA1002246	16095	16096
	Y79AA1002298	C-Y79AA1002298	16097	16098
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10	Y79AA1001145	C-Y79AA1001145	16015	16016
	Y79AA1001177	C-Y79AA1001177	16017	16018
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40	Y79AA1001665	C-Y79AA1001665	16056	
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50	Y79AA1001827	C-Y79AA1001827	16069	16070
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	Y79AA1000202	C-Y79AA1000202	15942	15943
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	Y79AA1000268	C-Y79AA1000268	15950	15951
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20	Y79AA1000328	C-Y79AA1000328	15954	15955
	Y79AA1000355	C-Y79AA1000355	15956	15957
	Y79AA1000368	C-Y79AA1000368	15958	15959
	Y79AA1000420	C-Y79AA1000420	15960	15961
25	Y79AA1000469	C-Y79AA1000469	15962	15963
	Y79AA1000480	C-Y79AA1000480	15964	15965
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	Y79AA1000705	C-Y79AA1000705	15974	15975
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	Y79AA1000752	C-Y79AA1000752	15980	15981
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	Y79AA1000794	C-Y79AA1000794	15988	15989
45	Y79AA1000800	C-Y79AA1000800	15990	15991
	Y79AA1000805	C-Y79AA1000805	15992	
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	Y79AA1000833	C-Y79AA1000833	15995	15996
50	Y79AA1000850	C-Y79AA1000850	15997	15998
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55	Y79AA1000976	C-Y79AA1000976	16003	16004

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25	THYR01001321	C-THYR01001321	15892	
	THYR01001322	C-THYR01001322	15893	15894
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	THYR01001656	C-THYR01001656	15912	15913
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	THYR01001809	C-THYR01001809	15926	15927
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	PLACE1011133	C-PLACE1011133	15674	
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	PLACE1011452	C-PLACE1011452	15694	
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	PLACE1002655	C-PLACE1002655	15554	15555
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	NT2RP3004258	C-NT2RP3004258	15215	15216
	NT2RP3004262	C-NT2RP3004262	15217	15218
15	NT2RP3004341	C-NT2RP3004341	15219	
	NT2RP3004378	C-NT2RP3004378	15220	15221
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	NT2RP3004428	C-NT2RP3004428	15224	15225
20	NT2RP3004451	C-NT2RP3004451	15226	15227
	NT2RP3004454	C-NT2RP3004454	15228	15229
	NT2RP3004472	C-NT2RP3004472	15230	15231
	NT2RP3004498	C-NT2RP3004498	15232	15233
25	NT2RP3004504	C-NT2RP3004504	15234	15235
	NT2RP3004507	C-NT2RP3004507	15236	15237
	NT2RP3004534	C-NT2RP3004534	15238	15239
30	NT2RP4000528	C-NT2RP4000528	15240	15241
	NT2RP4000907	C-NT2RP4000907	15242	15243
	NT2RP4001029	C-NT2RP4001029	15244	15245
	NT2RP4001336	C-NT2RP4001336	15246	15247
35	NT2RP4001389	C-NT2RP4001389	15248	15249
	NT2RP4001442	C-NT2RP4001442	15250	15251
	NT2RP4001529	C-NT2RP4001529	15252	15253
	NT2RP4001656	C-NT2RP4001656	15254	15255
40	OVARC1000106	C-OVARC1000106	15256	15257
	OVARC1000198	C-OVARC1000198	15258	
	OVARC1000682	C-OVARC1000682	15259	15260
45	OVARC1000703	C-OVARC1000703	15261	15262
	OVARC1000722	C-OVARC1000722	15263	15264
	OVARC1000730	C-OVARC1000730	15265	15266
	OVARC1000746	C-OVARC1000746	15267	15268
50	OVARC1000781	C-OVARC1000781	15269	15270
	OVARC1000787	C-OVARC1000787	15271	
	OVARC1000834	C-OVARC1000834	15272	15273
55	OVARC1000846	C-OVARC1000846	15274	15275

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	NT2RP3002146	C-NT2RP3002146	15130	15131
	NT2RP3002151	C-NT2RP3002151	15132	15133
5	NT2RP3002165	C-NT2RP3002165	15134	15135
	NT2RP3002166	C-NT2RP3002166	15136	15137
	NT2RP3002181	C-NT2RP3002181	15138	15139
	NT2RP3002244	C-NT2RP3002244	15140	15141
10	NT2RP3002248	C-NT2RP3002248	15142	15143
	NT2RP3002273	C-NT2RP3002273	15144	15145
	NT2RP3002276	C-NT2RP3002276	15146	
	NT2RP3002304	C-NT2RP3002304	15147	
15	NT2RP3002501	C-NT2RP3002501	15148	15149
	NT2RP3002529	C-NT2RP3002529	15150	15151
	NT2RP3002566	C-NT2RP3002566	15152	15153
20	NT2RP3002587	C-NT2RP3002587	15154	
	NT2RP3002590	C-NT2RP3002590	15155	15156
	NT2RP3002631	C-NT2RP3002631	15157	15158
	NT2RP3002650	C-NT2RP3002650	15159	15160
25	NT2RP3002663	C-NT2RP3002663	15161	15162
	NT2RP3002671	C-NT2RP3002671	15163	15164
	NT2RP3002763	C-NT2RP3002763	15165	15166
	NT2RP3002861	C-NT2RP3002861	15167	
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	NT2RP3002953	C-NT2RP3002953	15171	15172
35	NT2RP3002988	C-NT2RP3002988	15173	15174
	NT2RP3003008	C-NT2RP3003008	15175	15176
	NT2RP3003101	C-NT2RP3003101	15177	15178
	NT2RP3003204	C-NT2RP3003204	15179	15180
40	NT2RP3003278	C-NT2RP3003278	15181	15182
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	NT2RP3003290	C-NT2RP3003290	15185	15186
45	NT2RP3003302	C-NT2RP3003302	15187	15188
	NT2RP3003313	C-NT2RP3003313	15189	15190
	NT2RP3003327	C-NT2RP3003327	15191	
	NT2RP3003344	C-NT2RP3003344	15192	15193
50	NT2RP3003353	C-NT2RP3003353	15194	15195
	NT2RP3003377	C-NT2RP3003377	15196	15197
	NT2RP3003385	C-NT2RP3003385	15198	15199
55	NT2RP3003433	C-NT2RP3003433	15200	

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5	NT2RP3001449	C-NT2RP3001449	15062	15063
	NT2RP3001453	C-NT2RP3001453	15064	15065
	NT2RP3001459	C-NT2RP3001459	15066	
	NT2RP3001527	C-NT2RP3001527	15067	15068
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	NT2RP3001580	C-NT2RP3001580	15071	15072
	NT2RP3001587	C-NT2RP3001587	15073	15074
15	NT2RP3001589	C-NT2RP3001589	15075	15076
	NT2RP3001607	C-NT2RP3001607	15077	
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	NT2RP3001671	C-NT2RP3001671	15080	15081
20	NT2RP3001672	C-NT2RP3001672	15082	15083
	NT2RP3001678	C-NT2RP3001678	15084	15085
	NT2RP3001688	C-NT2RP3001688	15086	15087
	NT2RP3001690	C-NT2RP3001690	15088	15089
25	NT2RP3001698	C-NT2RP3001698	15090	
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	NT2RP3001716	C-NT2RP3001716	15093	15094
30	NT2RP3001752	C-NT2RP3001752	15095	
	NT2RP3001792	C-NT2RP3001792	15096	15097
	NT2RP3001844	C-NT2RP3001844	15098	
	NT2RP3001854	C-NT2RP3001854	15099	15100
35	NT2RP3001855	C-NT2RP3001855	15101	15102
	NT2RP3001898	C-NT2RP3001898	15103	15104
	NT2RP3001931	C-NT2RP3001931	15105	15106
	NT2RP3001969	C-NT2RP3001969	15107	15108
40	NT2RP3002002	C-NT2RP3002002	15109	15110
	NT2RP3002004	C-NT2RP3002004	15111	15112
	NT2RP3002007	C-NT2RP3002007	15113	15114
45	NT2RP3002014	C-NT2RP3002014	15115	15116
	NT2RP3002045	C-NT2RP3002045	15117	15118
	NT2RP3002056	C-NT2RP3002056	15119	15120
	NT2RP3002062	C-NT2RP3002062	15121	15122
50	NT2RP3002081	C-NT2RP3002081	15123	15124
	NT2RP3002097	C-NT2RP3002097	15125	15126
	NT2RP3002108	C-NT2RP3002108	15127	15128
55	NT2RP3002142	C-NT2RP3002142	15129	

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	NT2RP3000341	C-NT2RP3000341	14993	
	NT2RP3000393	C-NT2RP3000393	14994	14995
5	NT2RP3000441	C-NT2RP3000441	14996	14997
	NT2RP3000449	C-NT2RP3000449	14998	14999
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10	NT2RP3000456	C-NT2RP3000456	15002	15003
	NT2RP3000542	C-NT2RP3000542	15004	
	NT2RP3000561	C-NT2RP3000561	15005	
	NT2RP3000562	C-NT2RP3000562	15006	
15	NT2RP3000578	C-NT2RP3000578	15007	15008
	NT2RP3000590	C-NT2RP3000590	15009	15010
	NT2RP3000592	C-NT2RP3000592	15011	
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	NT2RP3000624	C-NT2RP3000624	15013	15014
	NT2RP3000685	C-NT2RP3000685	15015	15016
	NT2RP3000736	C-NT2RP3000736	15017	15018
25	NT2RP3000742	C-NT2RP3000742	15019	15020
	NT2RP3000753	C-NT2RP3000753	15021	15022
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	NT2RP3001007	C-NT2RP3001007	15028	
	NT2RP3001055	C-NT2RP3001055	15029	15030
35	NT2RP3001111	C-NT2RP3001111	15031	15032
	NT2RP3001120	C-NT2RP3001120	15033	15034
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40	NT2RP3001232	C-NT2RP3001232	15039	
	NT2RP3001268	C-NT2RP3001268	15040	15041
	NT2RP3001272	C-NT2RP3001272	15042	15043
	NT2RP3001274	C-NT2RP3001274	15044	15045
45	NT2RP3001281	C-NT2RP3001281	15046	15047
	NT2RP3001297	C-NT2RP3001297	15048	15049
	NT2RP3001318	C-NT2RP3001318	15050	
50	NT2RP3001338	C-NT2RP3001338	15051	15052
	NT2RP3001355	C-NT2RP3001355	15053	15054
	NT2RP3001374	C-NT2RP3001374	15055	15056
55	NT2RP3001428	C-NT2RP3001428	15057	15058

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	NT2RP2005038	C-NT2RP2005038	14929	14930
10	NT2RP2005162	C-NT2RP2005162	14931	14932
	NT2RP2005204	C-NT2RP2005204	14933	14934
	NT2RP2005227	C-NT2RP2005227	14935	14936
15	NT2RP2005287	C-NT2RP2005287	14937	
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	NT2RP2005539	C-NT2RP2005539	14942	14943
20	NT2RP2005605	C-NT2RP2005605	14944	14945
	NT2RP2005722	C-NT2RP2005722	14946	14947
	NT2RP2005732	C-NT2RP2005732	14948	14949
	NT2RP2005784	C-NT2RP2005784	14950	14951
25	NT2RP2005812	C-NT2RP2005812	14952	14953
	NT2RP2005859	C-NT2RP2005859	14954	14955
	NT2RP2006023	C-NT2RP2006023	14956	14957
30	NT2RP2006334	C-NT2RP2006334	14958	14959
	NT2RP2006441	C-NT2RP2006441	14960	
	NT2RP3000002	C-NT2RP3000002	14961	14962
	NT2RP3000050	C-NT2RP3000050	14963	14964
35	NT2RP3000055	C-NT2RP3000055	14965	14966
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40	NT2RP3000085	C-NT2RP3000085	14971	14972
	NT2RP3000092	C-NT2RP3000092	14973	14974
	NT2RP3000109	C-NT2RP3000109	14975	14976
	NT2RP3000134	C-NT2RP3000134	14977	14978
45	NT2RP3000149	C-NT2RP3000149	14979	14980
	NT2RP3000197	C-NT2RP3000197	14981	
	NT2RP3000207	C-NT2RP3000207	14982	14983
	NT2RP3000233	C-NT2RP3000233	14984	14985
50	NT2RP3000235	C-NT2RP3000235	14986	
	NT2RP3000247	C-NT2RP3000247	14987	
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55	NT2RP3000299	C-NT2RP3000299	14990	14991

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	NT2RP2003506	C-NT2RP2003506	14854	14855
	NT2RP2003511	C-NT2RP2003511	14856	14857
10	NT2RP2003513	C-NT2RP2003513	14858	14859
	NT2RP2003567	C-NT2RP2003567	14860	14861
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15	NT2RP2003691	C-NT2RP2003691	14864	
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	NT2RP2003764	C-NT2RP2003764	14869	14870
20	NT2RP2003769	C-NT2RP2003769	14871	
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	NT2RP2003840	C-NT2RP2003840	14874	14875
25	NT2RP2003857	C-NT2RP2003857	14876	14877
	NT2RP2003981	C-NT2RP2003981	14878	14879
	NT2RP2003984	C-NT2RP2003984	14880	14881
	NT2RP2004041	C-NT2RP2004041	14882	14883
30	NT2RP2004066	C-NT2RP2004066	14884	14885
	NT2RP2004081	C-NT2RP2004081	14886	14887
	NT2RP2004124	C-NT2RP2004124	14888	14889
	NT2RP2004152	C-NT2RP2004152	14890	14891
35	NT2RP2004165	C-NT2RP2004165	14892	14893
	NT2RP2004187	C-NT2RP2004187	14894	14895
	NT2RP2004239	C-NT2RP2004239	14896	14897
40	NT2RP2004245	C-NT2RP2004245	14898	14899
	NT2RP2004364	C-NT2RP2004364	14900	14901
	NT2RP2004365	C-NT2RP2004365	14902	14903
	NT2RP2004366	C-NT2RP2004366	14904	14905
45	NT2RP2004373	C-NT2RP2004373	14906	14907
	NT2RP2004476	C-NT2RP2004476	14908	14909
	NT2RP2004551	C-NT2RP2004551	14910	14911
50	NT2RP2004568	C-NT2RP2004568	14912	14913
	NT2RP2004600	C-NT2RP2004600	14914	14915
	NT2RP2004664	C-NT2RP2004664	14916	14917
	NT2RP2004743	C-NT2RP2004743	14918	
55	NT2RP2004768	C-NT2RP2004768	14919	14920

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5	NT2RP2002099	C-NT2RP2002099	14778	14779
	NT2RP2002185	C-NT2RP2002185	14780	14781
	NT2RP2002193	C-NT2RP2002193	14782	14783
10	NT2RP2002231	C-NT2RP2002231	14784	14785
	NT2RP2002235	C-NT2RP2002235	14786	14787
	NT2RP2002252	C-NT2RP2002252	14788	14789
	NT2RP2002292	C-NT2RP2002292	14790	14791
15	NT2RP2002408	C-NT2RP2002408	14792	14793
	NT2RP2002442	C-NT2RP2002442	14794	14795
	NT2RP2002464	C-NT2RP2002464	14796	14797
	NT2RP2002498	C-NT2RP2002498	14798	14799
20	NT2RP2002503	C-NT2RP2002503	14800	14801
	NT2RP2002520	C-NT2RP2002520	14802	14803
	NT2RP2002549	C-NT2RP2002549	14804	14805
25	NT2RP2002609	C-NT2RP2002609	14806	14807
	NT2RP2002706	C-NT2RP2002706	14808	
	NT2RP2002710	C-NT2RP2002710	14809	14810
	NT2RP2002800	C-NT2RP2002800	14811	
30	NT2RP2002880	C-NT2RP2002880	14812	14813
	NT2RP2002891	C-NT2RP2002891	14814	14815
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	NT2RP2002939	C-NT2RP2002939	14818	14819
35	NT2RP2002993	C-NT2RP2002993	14820	14821
	NT2RP2003034	C-NT2RP2003034	14822	
	NT2RP2003099	C-NT2RP2003099	14823	
40	NT2RP2003137	C-NT2RP2003137	14824	14825
	NT2RP2003157	C-NT2RP2003157	14826	14827
	NT2RP2003158	C-NT2RP2003158	14828	14829
	NT2RP2003165	C-NT2RP2003165	14830	
45	NT2RP2003243	C-NT2RP2003243	14831	14832
	NT2RP2003277	C-NT2RP2003277	14833	14834
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	NT2RP2003297	C-NT2RP2003297	14837	14838
50	NT2RP2003307	C-NT2RP2003307	14839	14840
	NT2RP2003308	C-NT2RP2003308	14841	14842
	NT2RP2003347	C-NT2RP2003347	14843	14844
55	NT2RP2003391	C-NT2RP2003391	14845	14846

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	NT2RP2001174	C-NT2RP2001174	14710	14711
	NT2RP2001196	C-NT2RP2001196	14712	
	NT2RP2001226	C-NT2RP2001226	14713	14714
10	NT2RP2001268	C-NT2RP2001268	14715	14716
	NT2RP2001290	C-NT2RP2001290	14717	14718
	NT2RP2001295	C-NT2RP2001295	14719	14720
	NT2RP2001312	C-NT2RP2001312	14721	
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	NT2RP2001328	C-NT2RP2001328	14724	
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	NT2RP2001392	C-NT2RP2001392	14729	14730
	NT2RP2001394	C-NT2RP2001394	14731	
	NT2RP2001420	C-NT2RP2001420	14732	14733
25	NT2RP2001450	C-NT2RP2001450	14734	
	NT2RP2001467	C-NT2RP2001467	14735	14736
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	NT2RP2001520	C-NT2RP2001520	14740	14741
	NT2RP2001536	C-NT2RP2001536	14742	14743
	NT2RP2001560	C-NT2RP2001560	14744	14745
35	NT2RP2001576	C-NT2RP2001576	14746	14747
	NT2RP2001581	C-NT2RP2001581	14748	14749
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	NT2RP2001628	C-NT2RP2001628	14752	14753
40	NT2RP2001663	C-NT2RP2001663	14754	
	NT2RP2001748	C-NT2RP2001748	14755	14756
	NT2RP2001813	C-NT2RP2001813	14757	
45	NT2RP2001883	C-NT2RP2001883	14758	14759
	NT2RP2001900	C-NT2RP2001900	14760	14761
	NT2RP2001947	C-NT2RP2001947	14762	14763
	NT2RP2001985	C-NT2RP2001985	14764	14765
50	NT2RP2001991	C-NT2RP2001991	14766	14767
	NT2RP2002025	C-NT2RP2002025	14768	14769
	NT2RP2002058	C-NT2RP2002058	14770	14771
55	NT2RP2002076	C-NT2RP2002076	14772	14773

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5	NT2RM4002623	C-NT2RM4002623	14635	14636
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	NT2RP1000363	C-NT2RP1000363	14639	14640
10	NT2RP1000418	C-NT2RP1000418	14641	14642
	NT2RP1000513	C-NT2RP1000513	14643	
	NT2RP1000721	C-NT2RP1000721	14644	14645
	NT2RP1000730	C-NT2RP1000730	14646	14647
15	NT2RP1000767	C-NT2RP1000767	14648	14649
	NT2RP1000836	C-NT2RP1000836	14650	14651
	NT2RP1000902	C-NT2RP1000902	14652	14653
	NT2RP1000943	C-NT2RP1000943	14654	14655
20	NT2RP1001033	C-NT2RP1001033	14656	14657
	NT2RP1001073	C-NT2RP1001073	14658	14659
	NT2RP1001199	C-NT2RP1001199	14660	14661
25	NT2RP1001248	C-NT2RP1001248	14662	14663
	NT2RP1001253	C-NT2RP1001253	14664	14665
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	NT2RP1001294	C-NT2RP1001294	14668	14669
30	NT2RP1001302	C-NT2RP1001302	14670	14671
	NT2RP1001310	C-NT2RP1001310	14672	14673
	NT2RP1001361	C-NT2RP1001361	14674	14675
35	NT2RP1001385	C-NT2RP1001385	14676	14677
	NT2RP1001432	C-NT2RP1001432	14678	
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	NT2RP2000076	C-NT2RP2000076	14681	14682
40	NT2RP2000098	C-NT2RP2000098	14683	14684
	NT2RP2000108	C-NT2RP2000108	14685	
	NT2RP2000257	C-NT2RP2000257	14686	14687
	NT2RP2000258	C-NT2RP2000258	14688	14689
45	NT2RP2000289	C-NT2RP2000289	14690	14691
	NT2RP2000327	C-NT2RP2000327	14692	
	NT2RP2000337	C-NT2RP2000337	14693	14694
50	NT2RP2000420	C-NT2RP2000420	14695	14696
	NT2RP2000459	C-NT2RP2000459	14697	14698
	NT2RP2000498	C-NT2RP2000498	14699	
	NT2RP2000758	C-NT2RP2000758	14700	14701
55	NT2RP2001137	C-NT2RP2001137	14702	14703

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	NT2RM4002034	C-NT2RM4002034	14558	14559
	NT2RM4002044	C-NT2RM4002044	14560	14561
5	NT2RM4002054	C-NT2RM4002054	14562	14563
	NT2RM4002063	C-NT2RM4002063	14564	14565
	NT2RM4002066	C-NT2RM4002066	14566	14567
	NT2RM4002075	C-NT2RM4002075	14568	14569
10	NT2RM4002128	C-NT2RM4002128	14570	14571
	NT2RM4002140	C-NT2RM4002140	14572	14573
	NT2RM4002145	C-NT2RM4002145	14574	14575
	NT2RM4002161	C-NT2RM4002161	14576	14577
15	NT2RM4002174	C-NT2RM4002174	14578	14579
	NT2RM4002189	C-NT2RM4002189	14580	14581
	NT2RM4002205	C-NT2RM4002205	14582	14583
20	NT2RM4002213	C-NT2RM4002213	14584	14585
	NT2RM4002226	C-NT2RM4002226	14586	14587
	NT2RM4002251	C-NT2RM4002251	14588	14589
	NT2RM4002256	C-NT2RM4002256	14590	14591
25	NT2RM4002266	C-NT2RM4002266	14592	14593
	NT2RM4002281	C-NT2RM4002281	14594	
	NT2RM4002287	C-NT2RM4002287	14595	14596
30	NT2RM4002294	C-NT2RM4002294	14597	14598
	NT2RM4002301	C-NT2RM4002301	14599	14600
	NT2RM4002323	C-NT2RM4002323	14601	14602
	NT2RM4002339	C-NT2RM4002339	14603	14604
35	NT2RM4002344	C-NT2RM4002344	14605	14606
	NT2RM4002373	C-NT2RM4002373	14607	14608
	NT2RM4002374	C-NT2RM4002374	14609	
	NT2RM4002383	C-NT2RM4002383	14610	
40	NT2RM4002409	C-NT2RM4002409	14611	14612
	NT2RM4002438	C-NT2RM4002438	14613	14614
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45	NT2RM4002452	C-NT2RM4002452	14617	14618
	NT2RM4002457	C-NT2RM4002457	14619	
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	NT2RM4002493	C-NT2RM4002493	14622	14623
50	NT2RM4002527	C-NT2RM4002527	14624	14625
	NT2RM4002532	C-NT2RM4002532	14626	14627
	NT2RM4002558	C-NT2RM4002558	14628	14629
55	NT2RM4002567	C-NT2RM4002567	14630	14631

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	NT2RM4001594	C-NT2RM4001594	14487	14488
	NT2RM4001597	C-NT2RM4001597	14489	14490
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	NT2RM4001629	C-NT2RM4001629	14493	14494
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	NT2RM4001682	C-NT2RM4001682	14501	14502
	NT2RM4001710	C-NT2RM4001710	14503	
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15	NT2RM4001715	C-NT2RM4001715	14506	14507
	NT2RM4001731	C-NT2RM4001731	14508	14509
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45	NT2RM4001953	C-NT2RM4001953	14544	
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50	NT2RM4001984	C-NT2RM4001984	14551	
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55	NT2RM4002018	C-NT2RM4002018	14556	14557

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	NT2RM4001320	C-NT2RM4001320	14443	14444
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	NT2RM4001384	C-NT2RM4001384	14455	14456
	NT2RM4001410	C-NT2RM4001410	14457	14458
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	NT2RM4001455	C-NT2RM4001455	14469	14470
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	NT2RM4001557	C-NT2RM4001557	14477	14478
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	NT2RM4001582	C-NT2RM4001582	14483	14484
55	NT2RM4001592	C-NT2RM4001592	14485	14486

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	NT2RM4000689	C-NT2RM4000689	14355	14356
15	NT2RM4000698	C-NT2RM4000698	14357	14358
	NT2RM4000700	C-NT2RM4000700	14359	14360
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	NT2RM4000717	C-NT2RM4000717	14363	14364
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	NT2RM4000741	C-NT2RM4000741	14369	14370
25	NT2RM4000751	C-NT2RM4000751	14371	14372
	NT2RM4000764	C-NT2RM4000764	14373	14374
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	NT2RM4000795	C-NT2RM4000795	14380	14381
	NT2RM4000796	C-NT2RM4000796	14382	14383
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	NT2RM4000813	C-NT2RM4000813	14386	14387
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40	NT2RM4000848	C-NT2RM4000848	14392	14393
	NT2RM4000852	C-NT2RM4000852	14394	
	NT2RM4000855	C-NT2RM4000855	14395	14396
	NT2RM4000887	C-NT2RM4000887	14397	14398
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	NT2RM4001054	C-NT2RM4001054	14409	14410
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	NT2RM2001700	C-NT2RM2001700	14270	14271
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	NT2RM2001768	C-NT2RM2001768	14284	14285
	NT2RM2001771	C-NT2RM2001771	14286	
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	NT2RM2001784	C-NT2RM2001784	14289	14290
	NT2RM2001785	C-NT2RM2001785	14291	14292
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	NT2RM2001879	C-NT2RM2001879	14304	
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30	NT2RM2002145	C-NT2RM2002145	14307	14308
	NT2RM4000027	C-NT2RM4000027	14309	14310
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	NT2RM4000202	C-NT2RM4000202	14325	14326
	NT2RM4000233	C-NT2RM4000233	14327	14328
45	NT2RM4000244	C-NT2RM4000244	14329	
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50	NT2RM4000327	C-NT2RM4000327	14335	
	NT2RM4000356	C-NT2RM4000356	14336	14337
	NT2RM4000425	C-NT2RM4000425	14338	
55	NT2RM4000433	C-NT2RM4000433	14339	14340

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	NT2RM1000874	C-NT2RM1000874	14194	14195
	NT2RM1000882	C-NT2RM1000882	14196	14197
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	NT2RM1000962	C-NT2RM1000962	14210	14211
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	NT2RM1001003	C-NT2RM1001003	14214	14215
	NT2RM1001043	C-NT2RM1001043	14216	
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	NT2RM1001085	C-NT2RM1001085	14221	14222
	NT2RM1001102	C-NT2RM1001102	14223	14224
	NT2RM1001105	C-NT2RM1001105	14225	14226
25	NT2RM1001139	C-NT2RM1001139	14227	14228
	NT2RM2000420	C-NT2RM2000420	14229	14230
	NT2RM2000566	C-NT2RM2000566	14231	14232
30	NT2RM2000609	C-NT2RM2000609	14233	14234
	NT2RM2000612	C-NT2RM2000612	14235	14236
	NT2RM2000735	C-NT2RM2000735	14237	14238
	NT2RM2001588	C-NT2RM2001588	14239	14240
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40	NT2RM2001652	C-NT2RM2001652	14249	14250
	NT2RM2001659	C-NT2RM2001659	14251	14252
	NT2RM2001664	C-NT2RM2001664	14253	14254
45	NT2RM2001668	C-NT2RM2001668	14255	14256
	NT2RM2001671	C-NT2RM2001671	14257	14258
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50	NT2RM2001688	C-NT2RM2001688	14263	14264
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	NT2RM2001696	C-NT2RM2001696	14266	14267
55	NT2RM2001698	C-NT2RM2001698	14268	14269

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	NT2RM1000186	C-NT2RM1000186	14122	14123
	NT2RM1000187	C-NT2RM1000187	14124	14125
5	NT2RM1000199	C-NT2RM1000199...	14126	14127
	NT2RM1000244	C-NT2RM1000244	14128	
	NT2RM1000252	C-NT2RM1000252	14129	14130
	NT2RM1000256	C-NT2RM1000256	14131	14132
10	NT2RM1000260	C-NT2RM1000260	14133	14134
	NT2RM1000271	C-NT2RM1000271	14135	14136
	NT2RM1000300	C-NT2RM1000300	14137	14138
	NT2RM1000314	C-NT2RM1000314	14139	14140
15	NT2RM1000354	C-NT2RM1000354	14141	14142
	NT2RM1000355	C-NT2RM1000355	14143	14144
	NT2RM1000365	C-NT2RM1000365	14145	
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	NT2RM1000388	C-NT2RM1000388	14148	14149
	NT2RM1000399	C-NT2RM1000399	14150	14151
	NT2RM1000430	C-NT2RM1000430	14152	14153
25	NT2RM1000555	C-NT2RM1000555	14154	14155
	NT2RM1000563	C-NT2RM1000563	14156	14157
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35	NT2RM1000699	C-NT2RM1000699	14167	
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	NT2RM1000746	C-NT2RM1000746	14171	14172
40	NT2RM1000770	C-NT2RM1000770	14173	14174
	NT2RM1000772	C-NT2RM1000772	14175	14176
	NT2RM1000780	C-NT2RM1000780	14177	14178
45	NT2RM1000800	C-NT2RM1000800	14179	14180
	NT2RM1000802	C-NT2RM1000802	14181	14182
	NT2RM1000811	C-NT2RM1000811	14183	14184
	NT2RM1000826	C-NT2RM1000826	14185	14186
50	NT2RM1000829	C-NT2RM1000829	14187	
	NT2RM1000850	C-NT2RM1000850	14188	14189
	NT2RM1000852	C-NT2RM1000852	14190	14191
55	NT2RM1000857	C-NT2RM1000857	14192	14193

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	MAMMA1003011	C-MAMMA1003011	14067	14068
15	MAMMA1003013	C-MAMMA1003013	14069	14070
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	MAMMA1003039	C-MAMMA1003039	14075	
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	MAMMA1003049	C-MAMMA1003049	14077	14078
25	MAMMA1003056	C-MAMMA1003056	14079	14080
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35	MAMMA1003135	C-MAMMA1003135	14090	14091
	MAMMA1003146	C-MAMMA1003146	14092	14093
	MAMMA1003150	C-MAMMA1003150	14094	14095
	MAMMA1003166	C-MAMMA1003166	14096	14097
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	NT2RM1000039	C-NT2RM1000039	14102	14103
45	NT2RM1000055	C-NT2RM1000055	14104	14105
	NT2RM1000059	C-NT2RM1000059	14106	14107
	NT2RM1000062	C-NT2RM1000062	14108	14109
	NT2RM1000118	C-NT2RM1000118	14110	14111
50	NT2RM1000119	C-NT2RM1000119	14112	14113
	NT2RM1000127	C-NT2RM1000127	14114	14115
	NT2RM1000131	C-NT2RM1000131	14116	14117
	NT2RM1000132	C-NT2RM1000132	14118	14119
55	NT2RM1000153	C-NT2RM1000153	14120	14121

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	MAMMA1002268	C-MAMMA1002268	13996	13997
	MAMMA1002269	C-MAMMA1002269	13998	
10	MAMMA1002292	C-MAMMA1002292	13999	14000
	MAMMA1002294	C-MAMMA1002294	14001	14002
	MAMMA1002297	C-MAMMA1002297	14003	14004
	MAMMA1002312	C-MAMMA1002312	14005	
15	MAMMA1002329	C-MAMMA1002329	14006	14007
	MAMMA1002333	C-MAMMA1002333	14008	14009
	MAMMA1002351	C-MAMMA1002351	14010	14011
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	MAMMA1002355	C-MAMMA1002355	14013	
	MAMMA1002356	C-MAMMA1002356	14014	
	MAMMA1002362	C-MAMMA1002362	14015	
25	MAMMA1002380	C-MAMMA1002380	14016	14017
	MAMMA1002384	C-MAMMA1002384	14018	
	MAMMA1002427	C-MAMMA1002427	14019	14020
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	MAMMA1002485	C-MAMMA1002485	14023	14024
	MAMMA1002494	C-MAMMA1002494	14025	
	MAMMA1002524	C-MAMMA1002524	14026	14027
35	MAMMA1002530	C-MAMMA1002530	14028	14029
	MAMMA1002554	C-MAMMA1002554	14030	
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	MAMMA1002598	C-MAMMA1002598	14033	14034
40	MAMMA1002619	C-MAMMA1002619	14035	14036
	MAMMA1002655	C-MAMMA1002655	14037	14038
	MAMMA1002671	C-MAMMA1002671	14039	14040
45	MAMMA1002673	C-MAMMA1002673	14041	
	MAMMA1002684	C-MAMMA1002684	14042	14043
	MAMMA1002711	C-MAMMA1002711	14044	
	MAMMA1002769	C-MAMMA1002769	14045	14046
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	HEMBA1001824	C-HEMBA1001824	13389	13390
	HEMBA1001866	C-HEMBA1001866	13391	13392
	HEMBA1001910	C-HEMBA1001910	13393	
10	HEMBA1001913	C-HEMBA1001913	13394	13395
	HEMBA1001921	C-HEMBA1001921	13396	13397
	HEMBA1001939	C-HEMBA1001939	13398	13399
	HEMBA1001950	C-HEMBA1001950	13400	13401
15	HEMBA1001967	C-HEMBA1001967	13402	13403
	HEMBA1002035	C-HEMBA1002035	13404	13405
	HEMBA1002092	C-HEMBA1002092	13406	13407
20	HEMBA1002102	C-HEMBA1002102	13408	13409
	HEMBA1002150	C-HEMBA1002150	13410	13411
	HEMBA1002151	C-HEMBA1002151	13412	13413
	HEMBA1002189	C-HEMBA1002189	13414	
25	HEMBA1002215	C-HEMBA1002215	13415	13416
	HEMBA1002229	C-HEMBA1002229	13417	
	HEMBA1002241	C-HEMBA1002241	13418	13419
30	HEMBA1002341	C-HEMBA1002341	13420	
	HEMBA1002417	C-HEMBA1002417	13421	13422
	HEMBA1002547	C-HEMBA1002547	13423	13424
	HEMBA1002703	C-HEMBA1002703	13425	13426
35	HEMBA1002779	C-HEMBA1002779	13427	
	HEMBA1002816	C-HEMBA1002816	13428	13429
	HEMBA1002970	C-HEMBA1002970	13430	
40	HEMBA1002999	C-HEMBA1002999	13431	13432
	HEMBA1003021	C-HEMBA1003021	13433	
	HEMBA1003077	C-HEMBA1003077	13434	13435
	HEMBA1003079	C-HEMBA1003079	13436	13437
45	HEMBA1003273	C-HEMBA1003273	13438	
	HEMBA1003304	C-HEMBA1003304	13439	
	HEMBA1003309	C-HEMBA1003309	13440	
	HEMBA1003376	C-HEMBA1003376	13441	
50	HEMBA1003384	C-HEMBA1003384	13442	
	HEMBA1003531	C-HEMBA1003531	13443	
	HEMBA1003548	C-HEMBA1003548	13444	
55	HEMBA1003556	C-HEMBA1003556	13445	13446

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	HEMBA1000523	C-HEMBA1000523	13328	13329
	HEMBA1000542	C-HEMBA1000542	13330	13331
5	HEMBA1000545	C-HEMBA1000545	13332	
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	HEMBA1000592	C-HEMBA1000592	13334	13335
	HEMBA1000594	C-HEMBA1000594	13336	
10	HEMBA1000604	C-HEMBA1000604	13337	
	HEMBA1000622	C-HEMBA1000622	13338	
	HEMBA1000637	C-HEMBA1000637	13339	13340
15	HEMBA1000655	C-HEMBA1000655	13341	
	HEMBA1000657	C-HEMBA1000657	13342	13343
	HEMBA1000749	C-HEMBA1000749	13344	
	HEMBA1000769	C-HEMBA1000769	13345	
20	HEMBA1000773	C-HEMBA1000773	13346	
	HEMBA1000774	C-HEMBA1000774	13347	
	HEMBA1000822	C-HEMBA1000822	13348	13349
25	HEMBA1000843	C-HEMBA1000843	13350	
	HEMBA1000852	C-HEMBA1000852	13351	
	HEMBA1000870	C-HEMBA1000870	13352	13353
	HEMBA1000908	C-HEMBA1000908	13354	13355
30	HEMBA1000934	C-HEMBA1000934	13356	
	HEMBA1000972	C-HEMBA1000972	13357	13358
	HEMBA1000986	C-HEMBA1000986	13359	
35	HEMBA1000991	C-HEMBA1000991	13360	13361
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	HEMBA1001059	C-HEMBA1001059	13363	13364
	HEMBA1001094	C-HEMBA1001094	13365	
40	HEMBA1001302	C-HEMBA1001302	13366	13367
	HEMBA1001330	C-HEMBA1001330	13368	
	HEMBA1001497	C-HEMBA1001497	13369	
	HEMBA1001569	C-HEMBA1001569	13370	
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	HEMBA1001620	C-HEMBA1001620	13372	13373
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	HEMBA1001672	C-HEMBA1001672	13376	13377
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55	HEMBA1001746	C-HEMBA1001746	13381	13382

Table 351

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HEMBA1000264	C-HEMBA1000264	13306	
HEMBA1000280	C-HEMBA1000280	13307	13308
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HEMBA1000504	C-HEMBA1000504	13324	
HEMBA1000518	C-HEMBA1000518	13325	13326
HEMBA1000519	C-HEMBA1000519	13327	

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	PLACE2000050	C-PLACE2000050	13279	
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	PLACE4000590	C-PLACE4000590	13282	13283
15	PLACE4000638	C-PLACE4000638	13284	13285
	PLACE4000650	C-PLACE4000650	13286	13287
20	Y79AA1001647	C-Y79AA1001647	13288	13289

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	PLACE1010031	C-PLACE1010031	13209	13210
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	PLACE1010096	C-PLACE1010096	13216	13217
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	PLACE1010105	C-PLACE1010105	13219	13220
	PLACE1010106	C-PLACE1010106	13221	
	PLACE1010134	C-PLACE1010134	13222	13223
15	PLACE1010148	C-PLACE1010148	13224	13225
	PLACE1010194	C-PLACE1010194	13226	13227
	PLACE1010202	C-PLACE1010202	13228	13229
20	PLACE1010261	C-PLACE1010261	13230	13231
	PLACE1010274	C-PLACE1010274	13232	13233
	PLACE1010293	C-PLACE1010293	13234	
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	PLACE1010491	C-PLACE1010491	13245	
35	PLACE1010492	C-PLACE1010492	13246	13247
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	PLACE1010547	C-PLACE1010547	13252	13253
40	PLACE1010599	C-PLACE1010599	13254	13255
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	PLACE1010661	C-PLACE1010661	13261	13262
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	PLACE1010771	C-PLACE1010771	13268	13269
55	PLACE1010786	C-PLACE1010786	13270	

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	PLACE1009388	C-PLACE1009388	13151	
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	PLACE1009476	C-PLACE1009476	13164	13165
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	PLACE1009571	C-PLACE1009571	13170	13171
	PLACE1009581	C-PLACE1009581	13172	13173
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40	PLACE1009708	C-PLACE1009708	13185	13186
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	PLACE1009886	C-PLACE1009886	13197	
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50	PLACE1009971	C-PLACE1009971	13200	
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	PLACE1009995	C-PLACE1009995	13203	13204
55	PLACE1009997	C-PLACE1009997	13205	13206

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	PLACE1008488	C-PLACE1008488	13074	13075
5	PLACE1008524	C-PLACE1008524	13076	
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	PLACE1008568	C-PLACE1008568	13083	13084
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	PLACE1008621	C-PLACE1008621	13087	
15	PLACE1008626	C-PLACE1008626	13088	
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25	PLACE1008808	C-PLACE1008808	13099	13100
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30	PLACE1008867	C-PLACE1008867	13105	13106
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35	PLACE1009020	C-PLACE1009020	13110	13111
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	PLACE1009060	C-PLACE1009060	13116	13117
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45	PLACE1009099	C-PLACE1009099	13122	13123
	PLACE1009110	C-PLACE1009110	13124	
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	PLACE1009166	C-PLACE1009166	13131	13132
	PLACE1009174	C-PLACE1009174	13133	13134
55	PLACE1009186	C-PLACE1009186	13135	13136

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	PLACE1008402	C-PLACE1008402	13067	13068
55	PLACE1008429	C-PLACE1008429	13069	13070

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	PLACE1007239	C-PLACE1007239	12938	12939
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	PLACE1007274	C-PLACE1007274	12945	12946
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	PLACE1007346	C-PLACE1007346	12953	12954
	PLACE1007367	C-PLACE1007367	12955	
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	PLACE1005955	C-PLACE1005955	12837	12838
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	PLACE1006119	C-PLACE1006119	12847	
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	PLACE1006159	C-PLACE1006159	12850	12851
50	PLACE1006167	C-PLACE1006167	12852	12853
	PLACE1006170	C-PLACE1006170	12854	12855
	PLACE1006195	C-PLACE1006195	12856	12857
55	PLACE1006196	C-PLACE1006196	12858	12859

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	PLACE1004672	C-PLACE1004672	12721	12722
	PLACE1004674	C-PLACE1004674	12723	12724
5	PLACE1004691	C-PLACE1004691	12725	
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	PLACE1004736	C-PLACE1004736	12728	12729
	PLACE1004740	C-PLACE1004740	12730	
10	PLACE1004743	C-PLACE1004743	12731	12732
	PLACE1004751	C-PLACE1004751	12733	12734
	PLACE1004777	C-PLACE1004777	12735	12736
15	PLACE1004804	C-PLACE1004804	12737	12738
	PLACE1004814	C-PLACE1004814	12739	12740
	PLACE1004824	C-PLACE1004824	12741	12742
	PLACE1004868	C-PLACE1004868	12743	12744
20	PLACE1004885	C-PLACE1004885	12745	12746
	PLACE1004902	C-PLACE1004902	12747	12748
	PLACE1004918	C-PLACE1004918	12749	12750
25	PLACE1004930	C-PLACE1004930	12751	12752
	PLACE1004934	C-PLACE1004934	12753	
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30	PLACE1004982	C-PLACE1004982	12758	12759
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	PLACE1005027	C-PLACE1005027	12762	
	PLACE1005046	C-PLACE1005046	12763	
35	PLACE1005077	C-PLACE1005077	12764	
	PLACE1005101	C-PLACE1005101	12765	12766
	PLACE1005102	C-PLACE1005102	12767	12768
40	PLACE1005111	C-PLACE1005111	12769	
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	PLACE1005187	C-PLACE1005187	12771	12772
	PLACE1005206	C-PLACE1005206	12773	12774
45	PLACE1005232	C-PLACE1005232	12775	
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	PLACE1005266	C-PLACE1005266	12780	
50	PLACE1005277	C-PLACE1005277	12781	12782
	PLACE1005287	C-PLACE1005287	12783	12784
	PLACE1005305	C-PLACE1005305	12785	12786
55	PLACE1005308	C-PLACE1005308	12787	12788

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	PLACE1003762	C-PLACE1003762	12651	12652
5	PLACE1003771	C-PLACE1003771	12653	12654
	PLACE1003784	C-PLACE1003784	12655	12656
	PLACE1003923	C-PLACE1003923	12657	12658
	PLACE1003936	C-PLACE1003936	12659	
10	PLACE1003968	C-PLACE1003968	12660	12661
	PLACE1004104	C-PLACE1004104	12662	12663
	PLACE1004114	C-PLACE1004114	12664	
	PLACE1004128	C-PLACE1004128	12665	12666
15	PLACE1004149	C-PLACE1004149	12667	12668
	PLACE1004156	C-PLACE1004156	12669	12670
	PLACE1004161	C-PLACE1004161	12671	12672
20	PLACE1004183	C-PLACE1004183	12673	12674
	PLACE1004197	C-PLACE1004197	12675	12676
	PLACE1004203	C-PLACE1004203	12677	12678
	PLACE1004258	C-PLACE1004258	12679	12680
25	PLACE1004270	C-PLACE1004270	12681	12682
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30	PLACE1004302	C-PLACE1004302	12687	12688
	PLACE1004316	C-PLACE1004316	12689	12690
	PLACE1004358	C-PLACE1004358	12691	12692
	PLACE1004376	C-PLACE1004376	12693	12694
35	PLACE1004388	C-PLACE1004388	12695	12696
	PLACE1004405	C-PLACE1004405	12697	
	PLACE1004428	C-PLACE1004428	12698	12699
	PLACE1004437	C-PLACE1004437	12700	12701
40	PLACE1004451	C-PLACE1004451	12702	
	PLACE1004460	C-PLACE1004460	12703	12704
	PLACE1004473	C-PLACE1004473	12705	12706
45	PLACE1004510	C-PLACE1004510	12707	12708
	PLACE1004516	C-PLACE1004516	12709	
	PLACE1004548	C-PLACE1004548	12710	
	PLACE1004564	C-PLACE1004564	12711	12712
50	PLACE1004629	C-PLACE1004629	12713	12714
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	PLACE1004646	C-PLACE1004646	12717	12718
55	PLACE1004664	C-PLACE1004664	12719	12720

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	PLACE1001928	C-PLACE1001928	12580	
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	PLACE1002073	C-PLACE1002073	12587	12588
	PLACE1002140	C-PLACE1002140	12589	12590
10	PLACE1002163	C-PLACE1002163	12591	
	PLACE1002170	C-PLACE1002170	12592	
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15	PLACE1002438	C-PLACE1002438	12595	12596
	PLACE1002465	C-PLACE1002465	12597	12598
	PLACE1002529	C-PLACE1002529	12599	12600
	PLACE1002685	C-PLACE1002685	12601	12602
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	PLACE1002794	C-PLACE1002794	12605	12606
	PLACE1002815	C-PLACE1002815	12607	
25	PLACE1002839	C-PLACE1002839	12608	
	PLACE1002851	C-PLACE1002851	12609	12610
	PLACE1002941	C-PLACE1002941	12611	12612
	PLACE1002996	C-PLACE1002996	12613	12614
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	PLACE1003092	C-PLACE1003092	12617	12618
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40	PLACE1003200	C-PLACE1003200	12627	
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	PLACE1003302	C-PLACE1003302	12630	12631
	PLACE1003334	C-PLACE1003334	12632	12633
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	PLACE1003353	C-PLACE1003353	12636	12637
	PLACE1003369	C-PLACE1003369	12638	12639
50	PLACE1003602	C-PLACE1003602	12640	12641
	PLACE1003611	C-PLACE1003611	12642	12643
	PLACE1003625	C-PLACE1003625	12644	12645
	PLACE1003704	C-PLACE1003704	12646	12647
55	PLACE1003711	C-PLACE1003711	12648	

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	PLACE1000420	C-PLACE1000420	12511	12512
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	PLACE1000562	C-PLACE1000562	12516	12517
	PLACE1000564	C-PLACE1000564	12518	12519
	PLACE1000588	C-PLACE1000588	12520	12521
10	PLACE1000596	C-PLACE1000596	12522	12523
	PLACE1000611	C-PLACE1000611	12524	12525
	PLACE1000636	C-PLACE1000636	12526	12527
	PLACE1000716	C-PLACE1000716	12528	
15	PLACE1000748	C-PLACE1000748	12529	12530
	PLACE1000755	C-PLACE1000755	12531	12532
	PLACE1000785	C-PLACE1000785	12533	
20	PLACE1000798	C-PLACE1000798	12534	
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	PLACE1000909	C-PLACE1000909	12537	12538
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25	PLACE1000972	C-PLACE1000972	12541	12542
	PLACE1000977	C-PLACE1000977	12543	12544
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30	PLACE1001092	C-PLACE1001092	12546	12547
	PLACE1001257	C-PLACE1001257	12548	12549
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	PLACE1001387	C-PLACE1001387	12552	12553
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40	PLACE1001503	C-PLACE1001503	12557	12558
	PLACE1001570	C-PLACE1001570	12559	12560
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	PLACE1001692	C-PLACE1001692	12563	12564
45	PLACE1001729	C-PLACE1001729	12565	12566
	PLACE1001739	C-PLACE1001739	12567	12568
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	PLACE1001810	C-PLACE1001810	12571	
50	PLACE1001817	C-PLACE1001817	12572	12573
	PLACE1001869	C-PLACE1001869	12574	12575
	PLACE1001912	C-PLACE1001912	12576	12577
55	PLACE1001920	C-PLACE1001920	12578	12579

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	OVARC1000437	C-OVARC1000437	12442	12443
5	OVARC1000443	C-OVARC1000443	12444	12445
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	OVARC1000465	C-OVARC1000465	12448	12449
	OVARC1000466	C-OVARC1000466	12450	12451
10	OVARC1000473	C-OVARC1000473	12452	12453
	OVARC1000479	C-OVARC1000479	12454	12455
	OVARC1000520	C-OVARC1000520	12456	12457
15	OVARC1000564	C-OVARC1000564	12458	12459
	OVARC1000576	C-OVARC1000576	12460	
	OVARC1000588	C-OVARC1000588	12461	12462
	OVARC1000605	C-OVARC1000605	12463	12464
20	OVARC1000640	C-OVARC1000640	12465	12466
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25	OVARC1000771	C-OVARC1000771	12471	12472
	OVARC1000959	C-OVARC1000959	12473	12474
	OVARC1001034	C-OVARC1001034	12475	12476
	OVARC1001038	C-OVARC1001038	12477	12478
30	OVARC1001065	C-OVARC1001065	12479	12480
	OVARC1001162	C-OVARC1001162	12481	
	OVARC1001243	C-OVARC1001243	12482	
35	OVARC1001296	C-OVARC1001296	12483	12484
	OVARC1001360	C-OVARC1001360	12485	
	OVARC1001381	C-OVARC1001381	12486	12487
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40	PLACE1000005	C-PLACE1000005	12490	
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	PLACE1000142	C-PLACE1000142	12493	12494
	PLACE1000184	C-PLACE1000184	12495	
45	PLACE1000185	C-PLACE1000185	12496	12497
	PLACE1000213	C-PLACE1000213	12498	12499
	PLACE1000347	C-PLACE1000347	12500	12501
50	PLACE1000374	C-PLACE1000374	12502	
	PLACE1000380	C-PLACE1000380	12503	12504
	PLACE1000383	C-PLACE1000383	12505	12506
	PLACE1000401	C-PLACE1000401	12507	12508
55	PLACE1000406	C-PLACE1000406	12509	12510

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	NT2RP4002081	C-NT2RP4002081	12368	12369
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	NT2RP4002408	C-NT2RP4002408	12372	12373
	NT2RP4002791	C-NT2RP4002791	12374	12375
	NT2RP4002888	C-NT2RP4002888	12376	12377
10	NT2RP4002905	C-NT2RP4002905	12378	12379
	NT2RP5003461	C-NT2RP5003461	12380	12381
	NT2RP5003477	C-NT2RP5003477	12382	12383
	NT2RP5003492	C-NT2RP5003492	12384	12385
15	NT2RP5003500	C-NT2RP5003500	12386	12387
	NT2RP5003506	C-NT2RP5003506	12388	12389
	NT2RP5003522	C-NT2RP5003522	12390	12391
20	NT2RP5003524	C-NT2RP5003524	12392	12393
	NT2RP5003534	C-NT2RP5003534	12394	12395
	OVARC1000006	C-OVARC1000006	12396	12397
	OVARC1000013	C-OVARC1000013	12398	12399
25	OVARC1000014	C-OVARC1000014	12400	12401
	OVARC1000035	C-OVARC1000035	12402	
	OVARC1000060	C-OVARC1000060	12403	12404
30	OVARC1000087	C-OVARC1000087	12405	12406
	OVARC1000091	C-OVARC1000091	12407	12408
	OVARC1000113	C-OVARC1000113	12409	
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35	OVARC1000148	C-OVARC1000148	12412	12413
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	OVARC1000212	C-OVARC1000212	12419	12420
	OVARC1000241	C-OVARC1000241	12421	12422
	OVARC1000288	C-OVARC1000288	12423	12424
45	OVARC1000304	C-OVARC1000304	12425	12426
	OVARC1000309	C-OVARC1000309	12427	12428
	OVARC1000321	C-OVARC1000321	12429	12430
	OVARC1000326	C-OVARC1000326	12431	12432
50	OVARC1000335	C-OVARC1000335	12433	12434
	OVARC1000347	C-OVARC1000347	12435	
	OVARC1000384	C-OVARC1000384	12436	12437
55	OVARC1000411	C-OVARC1000411	12438	12439

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	NT2RP4001592	C-NT2RP4001592	12297	12298
	NT2RP4001610	C-NT2RP4001610	12299	12300
10	NT2RP4001614	C-NT2RP4001614	12301	12302
	NT2RP4001634	C-NT2RP4001634	12303	12304
	NT2RP4001638	C-NT2RP4001638	12305	12306
15	NT2RP4001644	C-NT2RP4001644	12307	12308
	NT2RP4001677	C-NT2RP4001677	12309	12310
	NT2RP4001679	C-NT2RP4001679	12311	12312
	NT2RP4001696	C-NT2RP4001696	12313	12314
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	NT2RP4001838	C-NT2RP4001838	12335	12336
35	NT2RP4001861	C-NT2RP4001861	12337	12338
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	NT2RP4001896	C-NT2RP4001896	12341	12342
40	NT2RP4001901	C-NT2RP4001901	12343	12344
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	NT2RP4001938	C-NT2RP4001938	12347	12348
	NT2RP4001946	C-NT2RP4001946	12349	12350
45	NT2RP4001950	C-NT2RP4001950	12351	12352
	NT2RP4001953	C-NT2RP4001953	12353	
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50	NT2RP4001975	C-NT2RP4001975	12356	12357
	NT2RP4002018	C-NT2RP4002018	12358	12359
	NT2RP4002052	C-NT2RP4002052	12360	12361
	NT2RP4002058	C-NT2RP4002058	12362	12363
55	NT2RP4002071	C-NT2RP4002071	12364	12365

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	NT2RP4001150	C-NT2RP4001150	12218	12219
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	NT2RP4001210	C-NT2RP4001210	12226	12227
	NT2RP4001219	C-NT2RP4001219	12228	12229
	NT2RP4001228	C-NT2RP4001228	12230	12231
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20	NT2RP4001274	C-NT2RP4001274	12238	12239
	NT2RP4001276	C-NT2RP4001276	12240	12241
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	NT2RP4001373	C-NT2RP4001373	12258	12259
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	NT2RP4001474	C-NT2RP4001474	12270	12271
	NT2RP4001483	C-NT2RP4001483	12272	12273
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50	NT2RP4001547	C-NT2RP4001547	12281	12282
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	NT2RP4000739	C-NT2RP4000739	12141	12142
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	NT2RP4000817	C-NT2RP4000817	12145	12146
10	NT2RP4000833	C-NT2RP4000833	12147	12148
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	NT2RP4000839	C-NT2RP4000839	12151	12152
15	NT2RP4000855	C-NT2RP4000855	12153	12154
	NT2RP4000865	C-NT2RP4000865	12155	12156
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	NT2RP4000927	C-NT2RP4000927	12163	12164
	NT2RP4000928	C-NT2RP4000928	12165	12166
	NT2RP4000929	C-NT2RP4000929	12167	12168
25	NT2RP4000955	C-NT2RP4000955	12169	12170
	NT2RP4000973	C-NT2RP4000973	12171	12172
	NT2RP4000975	C-NT2RP4000975	12173	12174
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	NT2RP4000997	C-NT2RP4000997	12181	12182
35	NT2RP4001004	C-NT2RP4001004	12183	12184
	NT2RP4001006	C-NT2RP4001006	12185	12186
	NT2RP4001010	C-NT2RP4001010	12187	12188
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	NT2RP4001079	C-NT2RP4001079	12195	12196
45	NT2RP4001080	C-NT2RP4001080	12197	12198
	NT2RP4001086	C-NT2RP4001086	12199	12200
	NT2RP4001095	C-NT2RP4001095	12201	12202
	NT2RP4001100	C-NT2RP4001100	12203	12204
50	NT2RP4001117	C-NT2RP4001117	12205	12206
	NT2RP4001122	C-NT2RP4001122	12207	12208
	NT2RP4001126	C-NT2RP4001126	12209	12210
55	NT2RP4001138	C-NT2RP4001138	12211	12212

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	NT2RP4000212	C-NT2RP4000212	12064	12065
	NT2RP4000243	C-NT2RP4000243	12066	12067
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	NT2RP4000518	C-NT2RP4000518	12111	12112
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	NT2RP4000704	C-NT2RP4000704	12131	12132
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55	NT2RP3000759	C-NT2RP3000759	11911	11912

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	NT2RP2006237	C-NT2RP2006237	11837	11838
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	NT2RP2005001	C-NT2RP2005001	11707	11708
	NT2RP2005012	C-NT2RP2005012	11709	11710
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	NT2RP2005472	C-NT2RP2005472	11742	11743
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	NT2RP2003108	C-NT2RP2003108	11636	11637
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	NT2RP2003446	C-NT2RP2003446	11656	11657
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	NT2RP2004316	C-NT2RP2004316	11677	
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	NT2RP2004655	C-NT2RP2004655	11687	11688
	NT2RP2004689	C-NT2RP2004689	11689	11690
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	NT2RP2002154	C-NT2RP2002154	11596	11597
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	NT2RP2002385	C-NT2RP2002385	11606	11607
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	NT2RP2002621	C-NT2RP2002621	11617	11618
	NT2RP2002672	C-NT2RP2002672	11619	
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	NT2RP2000369	C-NT2RP2000369	11495	
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	NT2RP2000503	C-NT2RP2000503	11506	
	NT2RP2000510	C-NT2RP2000510	11507	11508
25	NT2RP2000516	C-NT2RP2000516	11509	11510
	NT2RP2000603	C-NT2RP2000603	11511	11512
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	NT2RP2000668	C-NT2RP2000668	11521	11522
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	NT2RP2000809	C-NT2RP2000809	11528	11529
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	NT2RP2000814	C-NT2RP2000814	11532	11533
	NT2RP2000816	C-NT2RP2000816	11534	11535
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	NT2RP2000863	C-NT2RP2000863	11541	11542
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	NT2RP2000892	C-NT2RP2000892	11545	11546
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	NT2RP2000932	C-NT2RP2000932	11549	11550
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	NT2RP1001543	C-NT2RP1001543	11409	11410
5	NT2RP1001546	C-NT2RP1001546	11411	11412
	NT2RP1001569	C-NT2RP1001569	11413	11414
	NT2RP1001616	C-NT2RP1001616	11415	
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	NT2RP2000006	C-NT2RP2000006	11418	
	NT2RP2000007	C-NT2RP2000007	11419	11420
	NT2RP2000008	C-NT2RP2000008	11421	11422
15	NT2RP2000032	C-NT2RP2000032	11423	11424
	NT2RP2000045	C-NT2RP2000045	11425	11426
	NT2RP2000054	C-NT2RP2000054	11427	11428
	NT2RP2000056	C-NT2RP2000056	11429	
20	NT2RP2000067	C-NT2RP2000067	11430	11431
	NT2RP2000070	C-NT2RP2000070	11432	11433
	NT2RP2000079	C-NT2RP2000079	11434	11435
25	NT2RP2000088	C-NT2RP2000088	11436	
	NT2RP2000091	C-NT2RP2000091	11437	11438
	NT2RP2000097	C-NT2RP2000097	11439	11440
	NT2RP2000114	C-NT2RP2000114	11441	11442
30	NT2RP2000120	C-NT2RP2000120	11443	11444
	NT2RP2000126	C-NT2RP2000126	11445	11446
	NT2RP2000133	C-NT2RP2000133	11447	11448
35	NT2RP2000147	C-NT2RP2000147	11449	11450
	NT2RP2000153	C-NT2RP2000153	11451	11452
	NT2RP2000157	C-NT2RP2000157	11453	11454
	NT2RP2000161	C-NT2RP2000161	11455	11456
40	NT2RP2000173	C-NT2RP2000173	11457	11458
	NT2RP2000175	C-NT2RP2000175	11459	11460
	NT2RP2000195	C-NT2RP2000195	11461	11462
45	NT2RP2000205	C-NT2RP2000205	11463	11464
	NT2RP2000208	C-NT2RP2000208	11465	11466
	NT2RP2000224	C-NT2RP2000224	11467	11468
	NT2RP2000232	C-NT2RP2000232	11469	
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	NT2RP2000239	C-NT2RP2000239	11472	11473
	NT2RP2000248	C-NT2RP2000248	11474	11475
	NT2RP2000270	C-NT2RP2000270	11476	11477
55	NT2RP2000274	C-NT2RP2000274	11478	11479

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	NT2RP1000416	C-NT2RP1000416	11333	11334
	NT2RP1000439	C-NT2RP1000439	11335	11336
5	NT2RP1000443	C-NT2RP1000443	11337	11338
	NT2RP1000470	C-NT2RP1000470	11339	11340
	NT2RP1000478	C-NT2RP1000478	11341	11342
10	NT2RP1000481	C-NT2RP1000481	11343	11344
	NT2RP1000493	C-NT2RP1000493	11345	11346
	NT2RP1000547	C-NT2RP1000547	11347	11348
	NT2RP1000574	C-NT2RP1000574	11349	11350
15	NT2RP1000581	C-NT2RP1000581	11351	
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	NT2RP1000688	C-NT2RP1000688	11354	11355
20	NT2RP1000695	C-NT2RP1000695	11356	11357
	NT2RP1000733	C-NT2RP1000733	11358	11359
	NT2RP1000738	C-NT2RP1000738	11360	11361
	NT2RP1000782	C-NT2RP1000782	11362	11363
25	NT2RP1000825	C-NT2RP1000825	11364	11365
	NT2RP1000833	C-NT2RP1000833	11366	11367
	NT2RP1000846	C-NT2RP1000846	11368	
	NT2RP1000851	C-NT2RP1000851	11369	
30	NT2RP1000856	C-NT2RP1000856	11370	11371
	NT2RP1000915	C-NT2RP1000915	11372	11373
	NT2RP1000947	C-NT2RP1000947	11374	11375
35	NT2RP1000954	C-NT2RP1000954	11376	11377
	NT2RP1000958	C-NT2RP1000958	11378	11379
	NT2RP1000959	C-NT2RP1000959	11380	11381
	NT2RP1000966	C-NT2RP1000966	11382	11383
40	NT2RP1000980	C-NT2RP1000980	11384	11385
	NT2RP1000988	C-NT2RP1000988	11386	11387
	NT2RP1001011	C-NT2RP1001011	11388	11389
	NT2RP1001014	C-NT2RP1001014	11390	11391
45	NT2RP1001395	C-NT2RP1001395	11392	11393
	NT2RP1001410	C-NT2RP1001410	11394	11395
	NT2RP1001424	C-NT2RP1001424	11396	
50	NT2RP1001449	C-NT2RP1001449	11397	11398
	NT2RP1001457	C-NT2RP1001457	11399	11400
	NT2RP1001466	C-NT2RP1001466	11401	11402
	NT2RP1001475	C-NT2RP1001475	11403	11404
55	NT2RP1001482	C-NT2RP1001482	11405	11406

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	NT2RM4000290	C-NT2RM4000290	11257	11258
	NT2RM4000344	C-NT2RM4000344	11259	11260
5	NT2RM4000349	C-NT2RM4000349	11261	11262
	NT2RM4000354	C-NT2RM4000354	11263	11264
	NT2RM4000386	C-NT2RM4000386	11265	11266
10	NT2RM4000395	C-NT2RM4000395	11267	11268
	NT2RM4000421	C-NT2RM4000421	11269	11270
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	NT2RM4000471	C-NT2RM4000471	11273	11274
15	NT2RM4000486	C-NT2RM4000486	11275	11276
	NT2RM4000496	C-NT2RM4000496	11277	11278
	NT2RM4000511	C-NT2RM4000511	11279	11280
	NT2RM4000515	C-NT2RM4000515	11281	11282
20	NT2RM4000520	C-NT2RM4000520	11283	11284
	NT2RM4000585	C-NT2RM4000585	11285	11286
	NT2RM4000595	C-NT2RM4000595	11287	11288
25	NT2RP1000018	C-NT2RP1000018	11289	11290
	NT2RP1000035	C-NT2RP1000035	11291	11292
	NT2RP1000040	C-NT2RP1000040	11293	11294
	NT2RP1000063	C-NT2RP1000063	11295	11296
30	NT2RP1000086	C-NT2RP1000086	11297	11298
	NT2RP1000101	C-NT2RP1000101	11299	11300
	NT2RP1000111	C-NT2RP1000111	11301	11302
35	NT2RP1000112	C-NT2RP1000112	11303	
	NT2RP1000124	C-NT2RP1000124	11304	11305
	NT2RP1000130	C-NT2RP1000130	11306	11307
	NT2RP1000163	C-NT2RP1000163	11308	11309
40	NT2RP1000170	C-NT2RP1000170	11310	11311
	NT2RP1000191	C-NT2RP1000191	11312	11313
	NT2RP1000202	C-NT2RP1000202	11314	11315
	NT2RP1000243	C-NT2RP1000243	11316	11317
45	NT2RP1000259	C-NT2RP1000259	11318	
	NT2RP1000272	C-NT2RP1000272	11319	11320
	NT2RP1000326	C-NT2RP1000326	11321	11322
50	NT2RP1000333	C-NT2RP1000333	11323	11324
	NT2RP1000348	C-NT2RP1000348	11325	11326
	NT2RP1000357	C-NT2RP1000357	11327	11328
	NT2RP1000376	C-NT2RP1000376	11329	11330
55	NT2RP1000413	C-NT2RP1000413	11331	11332

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	NT2RM2001499	C-NT2RM2001499	11181	11182
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5	NT2RM2001524	C-NT2RM2001524	11185	11186
	NT2RM2001544	C-NT2RM2001544	11187	11188
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	NT2RM2001575	C-NT2RM2001575	11191	11192
10	NT2RM2001582	C-NT2RM2001582	11193	11194
	NT2RM2001886	C-NT2RM2001886	11195	11196
	NT2RM2001896	C-NT2RM2001896	11197	11198
	NT2RM2001903	C-NT2RM2001903	11199	11200
15	NT2RM2001930	C-NT2RM2001930	11201	11202
	NT2RM2001935	C-NT2RM2001935	11203	11204
	NT2RM2001936	C-NT2RM2001936	11205	11206
20	NT2RM2001950	C-NT2RM2001950	11207	11208
	NT2RM2001982	C-NT2RM2001982	11209	11210
	NT2RM2001989	C-NT2RM2001989	11211	11212
	NT2RM2001997	C-NT2RM2001997	11213	11214
25	NT2RM2001998	C-NT2RM2001998	11215	11216
	NT2RM2002004	C-NT2RM2002004	11217	11218
	NT2RM2002014	C-NT2RM2002014	11219	11220
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35	NT2RM2002091	C-NT2RM2002091	11229	11230
	NT2RM2002100	C-NT2RM2002100	11231	11232
	NT2RM2002109	C-NT2RM2002109	11233	11234
40	NT2RM2002128	C-NT2RM2002128	11235	11236
	NT2RM2002142	C-NT2RM2002142	11237	11238
	NT2RM2002178	C-NT2RM2002178	11239	11240
	NT2RM4000024	C-NT2RM4000024	11241	11242
45	NT2RM4000061	C-NT2RM4000061	11243	
	NT2RM4000104	C-NT2RM4000104	11244	11245
	NT2RM4000139	C-NT2RM4000139	11246	11247
	NT2RM4000169	C-NT2RM4000169	11248	11249
50	NT2RM4000191	C-NT2RM4000191	11250	11251
	NT2RM4000197	C-NT2RM4000197	11252	
	NT2RM4000210	C-NT2RM4000210	11253	11254
55	NT2RM4000229	C-NT2RM4000229	11255	11256

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	NT2RM2000639	C-NT2RM2000639	11108	11109
	NT2RM2000649	C-NT2RM2000649	11110	11111
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	NT2RM2000691	C-NT2RM2000691	11114	11115
	NT2RM2000714	C-NT2RM2000714	11116	11117
10	NT2RM2000718	C-NT2RM2000718	11118	11119
	NT2RM2000740	C-NT2RM2000740	11120	11121
	NT2RM2000795	C-NT2RM2000795	11122	
	NT2RM2000821	C-NT2RM2000821	11123	11124
15	NT2RM2000837	C-NT2RM2000837	11125	11126
	NT2RM2000951	C-NT2RM2000951	11127	11128
	NT2RM2000952	C-NT2RM2000952	11129	11130
	NT2RM2000984	C-NT2RM2000984	11131	11132
20	NT2RM2001004	C-NT2RM2001004	11133	11134
	NT2RM2001035	C-NT2RM2001035	11135	11136
	NT2RM2001065	C-NT2RM2001065	11137	11138
25	NT2RM2001100	C-NT2RM2001100	11139	11140
	NT2RM2001131	C-NT2RM2001131	11141	11142
	NT2RM2001141	C-NT2RM2001141	11143	11144
	NT2RM2001152	C-NT2RM2001152	11145	11146
30	NT2RM2001177	C-NT2RM2001177	11147	
	NT2RM2001194	C-NT2RM2001194	11148	11149
	NT2RM2001196	C-NT2RM2001196	11150	11151
35	NT2RM2001201	C-NT2RM2001201	11152	11153
	NT2RM2001221	C-NT2RM2001221	11154	11155
	NT2RM2001238	C-NT2RM2001238	11156	11157
	NT2RM2001243	C-NT2RM2001243	11158	11159
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	NT2RM2001256	C-NT2RM2001256	11162	11163
	NT2RM2001291	C-NT2RM2001291	11164	
	NT2RM2001306	C-NT2RM2001306	11165	11166
45	NT2RM2001312	C-NT2RM2001312	11167	
	NT2RM2001319	C-NT2RM2001319	11168	11169
	NT2RM2001324	C-NT2RM2001324	11170	11171
50	NT2RM2001345	C-NT2RM2001345	11172	11173
	NT2RM2001370	C-NT2RM2001370	11174	11175
	NT2RM2001393	C-NT2RM2001393	11176	11177
	NT2RM2001420	C-NT2RM2001420	11178	
55	NT2RM2001424	C-NT2RM2001424	11179	11180

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	NT2RM2000032	C-NT2RM2000032	11034	
	NT2RM2000042	C-NT2RM2000042	11035	
5	NT2RM2000092	C-NT2RM2000092	11036	11037
	NT2RM2000093	C-NT2RM2000093	11038	11039
	NT2RM2000101	C-NT2RM2000101	11040	11041
10	NT2RM2000191	C-NT2RM2000191	11042	11043
	NT2RM2000192	C-NT2RM2000192	11044	11045
	NT2RM2000239	C-NT2RM2000239	11046	11047
	NT2RM2000250	C-NT2RM2000250	11048	11049
15	NT2RM2000259	C-NT2RM2000259	11050	11051
	NT2RM2000260	C-NT2RM2000260	11052	11053
	NT2RM2000287	C-NT2RM2000287	11054	11055
20	NT2RM2000322	C-NT2RM2000322	11056	11057
	NT2RM2000359	C-NT2RM2000359	11058	11059
	NT2RM2000363	C-NT2RM2000363	11060	11061
	NT2RM2000368	C-NT2RM2000368	11062	11063
25	NT2RM2000371	C-NT2RM2000371	11064	11065
	NT2RM2000374	C-NT2RM2000374	11066	11067
	NT2RM2000395	C-NT2RM2000395	11068	11069
30	NT2RM2000402	C-NT2RM2000402	11070	11071
	NT2RM2000407	C-NT2RM2000407	11072	11073
	NT2RM2000422	C-NT2RM2000422	11074	11075
	NT2RM2000452	C-NT2RM2000452	11076	11077
35	NT2RM2000469	C-NT2RM2000469	11078	11079
	NT2RM2000490	C-NT2RM2000490	11080	11081
	NT2RM2000502	C-NT2RM2000502	11082	11083
	NT2RM2000504	C-NT2RM2000504	11084	11085
40	NT2RM2000522	C-NT2RM2000522	11086	11087
	NT2RM2000540	C-NT2RM2000540	11088	11089
	NT2RM2000567	C-NT2RM2000567	11090	11091
45	NT2RM2000569	C-NT2RM2000569	11092	
	NT2RM2000577	C-NT2RM2000577	11093	11094
	NT2RM2000581	C-NT2RM2000581	11095	11096
	NT2RM2000588	C-NT2RM2000588	11097	11098
50	NT2RM2000594	C-NT2RM2000594	11099	11100
	NT2RM2000599	C-NT2RM2000599	11101	11102
	NT2RM2000624	C-NT2RM2000624	11103	11104
	NT2RM2000635	C-NT2RM2000635	11105	11106
55	NT2RM2000636	C-NT2RM2000636	11107	

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	HEMBB1001105	C-HEMBB1001105	10967	
	HEMBB1001117	C-HEMBB1001117	10968	
5	HEMBB1001126	C-HEMBB1001126	10969	10970
	HEMBB1001137	C-HEMBB1001137	10971	10972
	HEMBB1001151	C-HEMBB1001151	10973	10974
10	HEMBB1001153	C-HEMBB1001153	10975	10976
	HEMBB1001169	C-HEMBB1001169	10977	
	HEMBB1001175	C-HEMBB1001175	10978	10979
	HEMBB1001182	C-HEMBB1001182	10980	10981
15	HEMBB1001199	C-HEMBB1001199	10982	
	HEMBB1001210	C-HEMBB1001210	10983	
	HEMBB1001242	C-HEMBB1001242	10984	10985
	HEMBB1001288	C-HEMBB1001288	10986	10987
20	HEMBB1001289	C-HEMBB1001289	10988	10989
	HEMBB1001294	C-HEMBB1001294	10990	10991
	HEMBB1001314	C-HEMBB1001314	10992	10993
25	HEMBB1001331	C-HEMBB1001331	10994	10995
	HEMBB1001339	C-HEMBB1001339	10996	10997
	HEMBB1001346	C-HEMBB1001346	10998	10999
	HEMBB1001369	C-HEMBB1001369	11000	
30	HEMBB1001384	C-HEMBB1001384	11001	11002
	HEMBB1001387	C-HEMBB1001387	11003	11004
	MAMMA1002317	C-MAMMA1002317	11005	
35	MAMMA1002319	C-MAMMA1002319	11006	
	MAMMA1002385	C-MAMMA1002385	11007	11008
	NT2RM1000080	C-NT2RM1000080	11009	11010
	NT2RM1000242	C-NT2RM1000242	11011	
40	NT2RM1000257	C-NT2RM1000257	11012	11013
	NT2RM1000280	C-NT2RM1000280	11014	11015
	NT2RM1000669	C-NT2RM1000669	11016	11017
	NT2RM1000781	C-NT2RM1000781	11018	
45	NT2RM1000867	C-NT2RM1000867	11019	11020
	NT2RM1001008	C-NT2RM1001008	11021	11022
	NT2RM1001044	C-NT2RM1001044	11023	11024
50	NT2RM1001074	C-NT2RM1001074	11025	11026
	NT2RM1001115	C-NT2RM1001115	11027	11028
	NT2RM2000006	C-NT2RM2000006	11029	
	NT2RM2000013	C-NT2RM2000013	11030	11031
55	NT2RM2000030	C-NT2RM2000030	11032	11033

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5	HEMBB1000391	C-HEMBB1000391	10910	10911
	HEMBB1000438	C-HEMBB1000438	10912	
	HEMBB1000449	C-HEMBB1000449	10913	
10	HEMBB1000589	C-HEMBB1000589	10914	
	HEMBB1000591	C-HEMBB1000591	10915	
	HEMBB1000623	C-HEMBB1000623	10916	10917
	HEMBB1000630	C-HEMBB1000630	10918	10919
15	HEMBB1000631	C-HEMBB1000631	10920	10921
	HEMBB1000632	C-HEMBB1000632	10922	10923
	HEMBB1000671	C-HEMBB1000671	10924	
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	HEMBB1000705	C-HEMBB1000705	10927	
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	HEMBB1000725	C-HEMBB1000725	10929	10930
25	HEMBB1000763	C-HEMBB1000763	10931	10932
	HEMBB1000781	C-HEMBB1000781	10933	10934
	HEMBB1000789	C-HEMBB1000789	10935	10936
	HEMBB1000807	C-HEMBB1000807	10937	10938
30	HEMBB1000810	C-HEMBB1000810	10939	
	HEMBB1000848	C-HEMBB1000848	10940	10941
	HEMBB1000852	C-HEMBB1000852	10942	
35	HEMBB1000870	C-HEMBB1000870	10943	
	HEMBB1000887	C-HEMBB1000887	10944	
	HEMBB1000908	C-HEMBB1000908	10945	
	HEMBB1000927	C-HEMBB1000927	10946	10947
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	HEMBB1000973	C-HEMBB1000973	10950	10951
	HEMBB1000975	C-HEMBB1000975	10952	10953
	HEMBB1000985	C-HEMBB1000985	10954	10955
45	HEMBB1000991	C-HEMBB1000991	10956	
	HEMBB1001011	C-HEMBB1001011	10957	
	HEMBB1001014	C-HEMBB1001014	10958	
50	HEMBB1001024	C-HEMBB1001024	10959	
	HEMBB1001056	C-HEMBB1001056	10960	10961
	HEMBB1001058	C-HEMBB1001058	10962	10963
	HEMBB1001068	C-HEMBB1001068	10964	10965
55	HEMBB1001096	C-HEMBB1001096	10966	

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	HEMBA1004806	C-HEMBA1004806	10844	
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5	HEMBA1004850	C-HEMBA1004850	10847	10848
	HEMBA1004863	C-HEMBA1004863	10849	
	HEMBA1004923	C-HEMBA1004923	10850	10851
	HEMBA1004929	C-HEMBA1004929	10852	10853
10	HEMBA1004930	C-HEMBA1004930	10854	10855
	HEMBA1004933	C-HEMBA1004933	10856	10857
	HEMBA1004954	C-HEMBA1004954	10858	
15	HEMBA1004972	C-HEMBA1004972	10859	10860
	HEMBA1005475	C-HEMBA1005475	10861	
	HEMBA1005581	C-HEMBA1005581	10862	10863
	HEMBA1006248	C-HEMBA1006248	10864	10865
20	HEMBA1006310	C-HEMBA1006310	10866	10867
	HEMBA1006344	C-HEMBA1006344	10868	10869
	HEMBA1006377	C-HEMBA1006377	10870	10871
25	HEMBA1006467	C-HEMBA1006467	10872	
	HEMBA1006474	C-HEMBA1006474	10873	10874
	HEMBA1006530	C-HEMBA1006530	10875	
	HEMBA1006737	C-HEMBA1006737	10876	10877
30	HEMBA1006795	C-HEMBA1006795	10878	
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	HEMBA1006936	C-HEMBA1006936	10881	10882
35	HEMBA1007018	C-HEMBA1007018	10883	10884
	HEMBA1007342	C-HEMBA1007342	10885	
	HEMBA1000008	C-HEMBA1000008	10886	10887
	HEMBA1000018	C-HEMBA1000018	10888	
40	HEMBA1000024	C-HEMBA1000024	10889	
	HEMBA1000025	C-HEMBA1000025	10890	
	HEMBA1000036	C-HEMBA1000036	10891	10892
	HEMBA1000037	C-HEMBA1000037	10893	10894
45	HEMBA1000083	C-HEMBA1000083	10895	10896
	HEMBA1000103	C-HEMBA1000103	10897	
	HEMBA1000119	C-HEMBA1000119	10898	10899
50	HEMBA1000136	C-HEMBA1000136	10900	
	HEMBA1000215	C-HEMBA1000215	10901	
	HEMBA1000226	C-HEMBA1000226	10902	10903
	HEMBA1000244	C-HEMBA1000244	10904	10905
55	HEMBA1000266	C-HEMBA1000266	10906	10907

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	HEMBA1003555	C-HEMBA1003555	10778	10779
	HEMBA1003560	C-HEMBA1003560	10780	
5	HEMBA1003568	C-HEMBA1003568	10781	10782
	HEMBA1003569	C-HEMBA1003569	10783	10784
	HEMBA1003581	C-HEMBA1003581	10785	
	HEMBA1003591	C-HEMBA1003591	10786	10787
10	HEMBA1003615	C-HEMBA1003615	10788	10789
	HEMBA1003617	C-HEMBA1003617	10790	10791
	HEMBA1003621	C-HEMBA1003621	10792	
15	HEMBA1003662	C-HEMBA1003662	10793	10794
	HEMBA1003690	C-HEMBA1003690	10795	10796
	HEMBA1003711	C-HEMBA1003711	10797	10798
	HEMBA1003807	C-HEMBA1003807	10799	
20	HEMBA1003864	C-HEMBA1003864	10800	10801
	HEMBA1003953	C-HEMBA1003953	10802	10803
	HEMBA1003959	C-HEMBA1003959	10804	
25	HEMBA1003989	C-HEMBA1003989	10805	10806
	HEMBA1004074	C-HEMBA1004074	10807	
	HEMBA1004097	C-HEMBA1004097	10808	10809
	HEMBA1004146	C-HEMBA1004146	10810	10811
30	HEMBA1004199	C-HEMBA1004199	10812	10813
	HEMBA1004207	C-HEMBA1004207	10814	
	HEMBA1004227	C-HEMBA1004227	10815	10816
	HEMBA1004246	C-HEMBA1004246	10817	
35	HEMBA1004276	C-HEMBA1004276	10818	10819
	HEMBA1004289	C-HEMBA1004289	10820	10821
	HEMBA1004509	C-HEMBA1004509	10822	10823
40	HEMBA1004534	C-HEMBA1004534	10824	10825
	HEMBA1004596	C-HEMBA1004596	10826	10827
	HEMBA1004693	C-HEMBA1004693	10828	10829
	HEMBA1004736	C-HEMBA1004736	10830	
45	HEMBA1004753	C-HEMBA1004753	10831	
	HEMBA1004756	C-HEMBA1004756	10832	10833
	HEMBA1004758	C-HEMBA1004758	10834	10835
	HEMBA1004763	C-HEMBA1004763	10836	10837
50	HEMBA1004768	C-HEMBA1004768	10838	10839
	HEMBA1004771	C-HEMBA1004771	10840	
	HEMBA1004776	C-HEMBA1004776	10841	
55	HEMBA1004795	C-HEMBA1004795	10842	10843

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	HEMBA1002971	C-HEMBA1002971	10710	
	HEMBA1002973	C-HEMBA1002973	10711	10712
5	HEMBA1002997	C-HEMBA1002997	10713	10714
	HEMBA1003033	C-HEMBA1003033	10715	10716
	HEMBA1003035	C-HEMBA1003035	10717	
	HEMBA1003041	C-HEMBA1003041	10718	10719
10	HEMBA1003046	C-HEMBA1003046	10720	10721
	HEMBA1003067	C-HEMBA1003067	10722	
	HEMBA1003096	C-HEMBA1003096	10723	10724
15	HEMBA1003117	C-HEMBA1003117	10725	10726
	HEMBA1003129	C-HEMBA1003129	10727	
	HEMBA1003136	C-HEMBA1003136	10728	10729
	HEMBA1003148	C-HEMBA1003148	10730	10731
20	HEMBA1003175	C-HEMBA1003175	10732	10733
	HEMBA1003179	C-HEMBA1003179	10734	10735
	HEMBA1003199	C-HEMBA1003199	10736	10737
25	HEMBA1003222	C-HEMBA1003222	10738	10739
	HEMBA1003235	C-HEMBA1003235	10740	10741
	HEMBA1003250	C-HEMBA1003250	10742	10743
	HEMBA1003257	C-HEMBA1003257	10744	10745
30	HEMBA1003281	C-HEMBA1003281	10746	10747
	HEMBA1003286	C-HEMBA1003286	10748	10749
	HEMBA1003291	C-HEMBA1003291	10750	10751
	HEMBA1003322	C-HEMBA1003322	10752	
35	HEMBA1003327	C-HEMBA1003327	10753	
	HEMBA1003369	C-HEMBA1003369	10754	10755
	HEMBA1003370	C-HEMBA1003370	10756	
40	HEMBA1003380	C-HEMBA1003380	10757	
	HEMBA1003395	C-HEMBA1003395	10758	10759
	HEMBA1003402	C-HEMBA1003402	10760	
	HEMBA1003408	C-HEMBA1003408	10761	10762
45	HEMBA1003417	C-HEMBA1003417	10763	10764
	HEMBA1003418	C-HEMBA1003418	10765	10766
	HEMBA1003433	C-HEMBA1003433	10767	10768
50	HEMBA1003447	C-HEMBA1003447	10769	10770
	HEMBA1003461	C-HEMBA1003461	10771	10772
	HEMBA1003463	C-HEMBA1003463	10773	
	HEMBA1003528	C-HEMBA1003528	10774	10775
55	HEMBA1003545	C-HEMBA1003545	10776	10777

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	HEMBA1002212	C-HEMBA1002212	10644	10645
	HEMBA1002237	C-HEMBA1002237	10646	
5	HEMBA1002265	C-HEMBA1002265	10647	
	HEMBA1002267	C-HEMBA1002267	10648	
	HEMBA1002349	C-HEMBA1002349	10649	
10	HEMBA1002363	C-HEMBA1002363	10650	10651
	HEMBA1002419	C-HEMBA1002419	10652	10653
	HEMBA1002430	C-HEMBA1002430	10654	
	HEMBA1002439	C-HEMBA1002439	10655	
15	HEMBA1002458	C-HEMBA1002458	10656	10657
	HEMBA1002460	C-HEMBA1002460	10658	
	HEMBA1002462	C-HEMBA1002462	10659	10660
	HEMBA1002469	C-HEMBA1002469	10661	10662
20	HEMBA1002475	C-HEMBA1002475	10663	10664
	HEMBA1002477	C-HEMBA1002477	10665	10666
	HEMBA1002495	C-HEMBA1002495	10667	10668
25	HEMBA1002515	C-HEMBA1002515	10669	
	HEMBA1002542	C-HEMBA1002542	10670	
	HEMBA1002569	C-HEMBA1002569	10671	10672
	HEMBA1002583	C-HEMBA1002583	10673	
30	HEMBA1002609	C-HEMBA1002609	10674	10675
	HEMBA1002624	C-HEMBA1002624	10676	10677
	HEMBA1002688	C-HEMBA1002688	10678	10679
	HEMBA1002696	C-HEMBA1002696	10680	10681
35	HEMBA1002750	C-HEMBA1002750	10682	
	HEMBA1002768	C-HEMBA1002768	10683	10684
	HEMBA1002770	C-HEMBA1002770	10685	10686
40	HEMBA1002777	C-HEMBA1002777	10687	10688
	HEMBA1002794	C-HEMBA1002794	10689	10690
	HEMBA1002810	C-HEMBA1002810	10691	10692
	HEMBA1002818	C-HEMBA1002818	10693	10694
45	HEMBA1002850	C-HEMBA1002850	10695	
	HEMBA1002863	C-HEMBA1002863	10696	10697
	HEMBA1002876	C-HEMBA1002876	10698	10699
50	HEMBA1002935	C-HEMBA1002935	10700	10701
	HEMBA1002937	C-HEMBA1002937	10702	10703
	HEMBA1002939	C-HEMBA1002939	10704	10705
	HEMBA1002951	C-HEMBA1002951	10706	10707
55	HEMBA1002954	C-HEMBA1002954	10708	10709

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	HEMBA1001351	C-HEMBA1001351	10578	10579
	HEMBA1001387	C-HEMBA1001387	10580	10581
5	HEMBA1001388	C-HEMBA1001388	10582	10583
	HEMBA1001398	C-HEMBA1001398	10584	10585
	HEMBA1001405	C-HEMBA1001405	10586	10587
10	HEMBA1001407	C-HEMBA1001407	10588	10589
	HEMBA1001413	C-HEMBA1001413	10590	10591
	HEMBA1001415	C-HEMBA1001415	10592	10593
	HEMBA1001446	C-HEMBA1001446	10594	10595
15	HEMBA1001450	C-HEMBA1001450	10596	
	HEMBA1001455	C-HEMBA1001455	10597	10598
	HEMBA1001510	C-HEMBA1001510	10599	10600
20	HEMBA1001526	C-HEMBA1001526	10601	10602
	HEMBA1001533	C-HEMBA1001533	10603	
	HEMBA1001579	C-HEMBA1001579	10604	10605
	HEMBA1001581	C-HEMBA1001581	10606	
25	HEMBA1001595	C-HEMBA1001595	10607	10608
	HEMBA1001635	C-HEMBA1001635	10609	10610
	HEMBA1001661	C-HEMBA1001661	10611	10612
	HEMBA1001702	C-HEMBA1001702	10613	
30	HEMBA1001714	C-HEMBA1001714	10614	
	HEMBA1001731	C-HEMBA1001731	10615	
	HEMBA1001744	C-HEMBA1001744	10616	10617
35	HEMBA1001809	C-HEMBA1001809	10618	
	HEMBA1001815	C-HEMBA1001815	10619	
	HEMBA1001819	C-HEMBA1001819	10620	10621
	HEMBA1001847	C-HEMBA1001847	10622	10623
40	HEMBA1001864	C-HEMBA1001864	10624	
	HEMBA1001869	C-HEMBA1001869	10625	10626
	HEMBA1001896	C-HEMBA1001896	10627	10628
	HEMBA1001987	C-HEMBA1001987	10629	
45	HEMBA1002018	C-HEMBA1002018	10630	10631
	HEMBA1002049	C-HEMBA1002049	10632	
	HEMBA1002084	C-HEMBA1002084	10633	10634
50	HEMBA1002125	C-HEMBA1002125	10635	10636
	HEMBA1002161	C-HEMBA1002161	10637	10638
	HEMBA1002177	C-HEMBA1002177	10639	10640
	HEMBA1002191	C-HEMBA1002191	10641	
55	HEMBA1002199	C-HEMBA1002199	10642	10643

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	HEMBA1000636	C-HEMBA1000636	10514	10515
	HEMBA1000682	C-HEMBA1000682	10516	
5	HEMBA1000686	C-HEMBA1000686	10517	10518
	HEMBA1000719	C-HEMBA1000719	10519	10520
	HEMBA1000727	C-HEMBA1000727	10521	10522
10	HEMBA1000752	C-HEMBA1000752	10523	
	HEMBA1000817	C-HEMBA1000817	10524	10525
	HEMBA1000851	C-HEMBA1000851	10526	10527
	HEMBA1000867	C-HEMBA1000867	10528	
15	HEMBA1000869	C-HEMBA1000869	10529	
	HEMBA1000872	C-HEMBA1000872	10530	10531
	HEMBA1000910	C-HEMBA1000910	10532	10533
20	HEMBA1000918	C-HEMBA1000918	10534	
	HEMBA1000919	C-HEMBA1000919	10535	10536
	HEMBA1000946	C-HEMBA1000946	10537	10538
	HEMBA1000968	C-HEMBA1000968	10539	
25	HEMBA1000971	C-HEMBA1000971	10540	10541
	HEMBA1000975	C-HEMBA1000975	10542	
	HEMBA1001009	C-HEMBA1001009	10543	10544
30	HEMBA1001022	C-HEMBA1001022	10545	
	HEMBA1001043	C-HEMBA1001043	10546	10547
	HEMBA1001052	C-HEMBA1001052	10548	10549
	HEMBA1001080	C-HEMBA1001080	10550	
35	HEMBA1001085	C-HEMBA1001085	10551	10552
	HEMBA1001088	C-HEMBA1001088	10553	10554
	HEMBA1001109	C-HEMBA1001109	10555	
	HEMBA1001122	C-HEMBA1001122	10556	
40	HEMBA1001133	C-HEMBA1001133	10557	
	HEMBA1001137	C-HEMBA1001137	10558	10559
	HEMBA1001140	C-HEMBA1001140	10560	10561
45	HEMBA1001174	C-HEMBA1001174	10562	10563
	HEMBA1001197	C-HEMBA1001197	10564	10565
	HEMBA1001235	C-HEMBA1001235	10566	
	HEMBA1001257	C-HEMBA1001257	10567	10568
50	HEMBA1001281	C-HEMBA1001281	10569	10570
	HEMBA1001286	C-HEMBA1001286	10571	10572
	HEMBA1001303	C-HEMBA1001303	10573	
	HEMBA1001310	C-HEMBA1001310	10574	10575
55	HEMBA1001326	C-HEMBA1001326	10576	10577

Table 350

	clone name	name of full-length nucleotide sequence	SEQ ID of full-length nucleotide sequence	SEQ ID of deduced amino acid sequence
5				
10				
	HEMBA1000005	C-HEMBA1000005	10468	10469
	HEMBA1000030	C-HEMBA1000030	10470	
15	HEMBA1000046	C-HEMBA1000046	10471	
	HEMBA1000050	C-HEMBA1000050	10472	
	HEMBA1000076	C-HEMBA1000076	10473	10474
20	HEMBA1000156	C-HEMBA1000156	10475	10476
	HEMBA1000158	C-HEMBA1000158	10477	10478
	HEMBA1000168	C-HEMBA1000168	10479	10480
	HEMBA1000185	C-HEMBA1000185	10481	10482
25	HEMBA1000193	C-HEMBA1000193	10483	10484
	HEMBA1000227	C-HEMBA1000227	10485	10486
	HEMBA1000288	C-HEMBA1000288	10487	
30	HEMBA1000302	C-HEMBA1000302	10488	
	HEMBA1000304	C-HEMBA1000304	10489	10490
	HEMBA1000307	C-HEMBA1000307	10491	10492
35	HEMBA1000369	C-HEMBA1000369	10493	10494
	HEMBA1000387	C-HEMBA1000387	10495	
	HEMBA1000392	C-HEMBA1000392	10496	
	HEMBA1000460	C-HEMBA1000460	10497	
40	HEMBA1000488	C-HEMBA1000488	10498	10499
	HEMBA1000491	C-HEMBA1000491	10500	10501
	HEMBA1000501	C-HEMBA1000501	10502	
45	HEMBA1000508	C-HEMBA1000508	10503	
	HEMBA1000520	C-HEMBA1000520	10504	
	HEMBA1000531	C-HEMBA1000531	10505	10506
50	HEMBA1000534	C-HEMBA1000534	10507	
	HEMBA1000555	C-HEMBA1000555	10508	10509
	HEMBA1000568	C-HEMBA1000568	10510	
	HEMBA1000588	C-HEMBA1000588	10511	
55	HEMBA1000608	C-HEMBA1000608	10512	10513

Table 349

5	Y79AA1002361	5.46	3.35	2.57	6.5	7.83	6.14	2.75	4.60	4.6	*	+		
	Y79AA1002365	1.93	1.66	1.86	2.93	2.21	2.54	1.34	2.05	2.05	*	+		
	Y79AA1002373	3.38	1.43	1.37	3.37	3.29	2.38	2.95	2.21	2.21				
	Y79AA1002376	434.81	300.04	466.40	120.28	171.61	120.00	316.81	454.58	454.6	**	-		
	Y79AA1002378	5.45	6.92	5.32	7.99	10.13	8.03	4.87	4.92	4.92	*	+		
10	Y79AA1002381	11.63	11.08	9.56	16.28	16.98	14.53	7.89	7.01	7.01	**	+	**	-
	Y79AA1002388	4.34	4.47	7.01	11.41	12.79	9.45	5.70	6.37	6.37	*	+		
	Y79AA1002399	4.43	1.48	1.47	4.2	2.82	2.25	3.39	3.35	3.35				
	Y79AA1002407	1.81	1.09	1.32	2.36	2.58	2.43	1.55	2.35	2.35	**	+		
	Y79AA1002413	15.88	6.76	10.60	19.95	26.46	17.33	9.58	12.56	12.56				
15	Y79AA1002416	5.12	2.89	2.97	4.45	4.32	5.10	4.13	4.19	4.19				
	Y79AA1002429	2.82	1.17	1.77	2.75	1.85	2.91	4.10	5.62	5.62		*	+	
	Y79AA1002431	4.04	2.82	3.86	2.55	4.38	4.86	4.06	5.56	5.56	-			
	Y79AA1002433	11.76	5.78	6.28	9.49	4.53	7.78	4.34	8.17	8.17				
	Y79AA1002445	10.95	9.11	9.11	11.15	8.78	14.80	10.37	11.14	11.14				
20	Y79AA1002461	10.04	5.58	4.92	9.55	8.99	8.05	5.89	7.75	7.75				
	Y79AA1002466	22.18	13.94	11.33	23.59	18.02	25.25	10.79	17.76	17.76				
	Y79AA1002471	5.76	3.00	5.65	6.94	8.49	9.26	5.31	7.89	7.89	*	+		
	Y79AA1002472	12.12	5.83	9.20	16.86	14.60	20.34	6.74	12.38	12.38	*	+		
25	Y79AA1002474	3.46	0.84	1.92	1.74	1.49	1.64	2.77	1.35	1.35				
	Y79AA1002482	13.92	8.55	11.10	23.82	23.90	29.62	10.40	14.99	14.99	**	+		
	Y79AA1002487	1.72	0.87	1.11	1.3	1.59	1.75	1.57	1.93	1.93				
	Y79AA1002490	13.58	4.80	6.45	5.13	6.72	3.78	4.31	7.19	7.19				
	Y79AA1002493	5.77	2.96	3.11	8.04	10.37	7.90	4.77	5.75	5.75	*	+		
30	ZRV6C1006278	1.43	0.95	1.01	1.16	2.05	0.47	1.35	2.06	2.06				

[0245] The correspondence of the full-length nucleotide sequences of the present invention and the corresponding deduced amino acid sequences with the clone names are shown below.

Table 348

	Y79AA1002012	3.88	1.69	1.78	4.4	6.99	4.19	2.25	2.80	2.8			
5	Y79AA1002017	4.13	2.53	3.93	3.44	3.03	1.90	3.46	3.57	3.57			
	Y79AA1002022	14.79	9.29	9.45	11.91	10.49	14.24	13.65	16.25	16.25			
	Y79AA1002027	2.08	0.73	0.78	2.44	1.84	1.40	2.55	2.70	2.7		*	+
	Y79AA1002050	9.08	4.52	6.60	9.28	6.06	9.49	5.33	7.52	7.52			
	Y79AA1002058	11.36	5.78	6.33	12.51	9.30	13.02	7.69	9.93	9.93			
10	Y79AA1002060	25.88	13.74	19.34	20.14	18.93	22.49	14.01	18.58	18.58			
	Y79AA1002062	13.71	6.57	6.87	16.86	16.66	14.29	6.71	8.83	8.83	*	+	
	Y79AA1002065	12.17	6.23	5.09	7.95	5.75	3.68	6.63	7.77	7.77			
	Y79AA1002067	14.5	8.32	9.44	2.21	3.03	2.42	3.46	4.06	4.06	*	-	*
	Y79AA1002069	7.51	3.78	4.23	4.94	4.88	2.84	3.88	6.24	6.24			
15	Y79AA1002070	60.51	38.18	52.01	44.77	31.84	34.13	26.73	37.56	37.56			
	Y79AA1002074	151.4	80.88	106.02	132.97	122.53	136.83	70.79	85.36	85.36			
	Y79AA1002076	2.73	1.63	2.34	2.2	2.35	2.60	2.59	2.75	2.75			
	Y79AA1002083	5	2.28	2.46	3.91	2.83	3.75	3.56	3.71	3.71			
	Y79AA1002084	5.09	3.13	3.61	5.26	3.68	3.36	3.65	3.99	3.99			
20	Y79AA1002086	7.09	2.92	3.98	4.7	3.74	3.75	3.43	4.46	4.46			
	Y79AA1002087	17.27	8.44	10.83	14.51	15.32	11.91	7.90	9.56	9.56			
	Y79AA1002089	5.98	2.23	2.36	4.43	5.76	5.05	4.46	3.99	3.99			
	Y79AA1002093	4.42	1.41	2.73	3.3	2.91	3.64	2.40	3.24	3.24			
	Y79AA1002101	7.66	3.43	4.43	3.23	2.81	2.96	1.93	9.08	9.08			
25	Y79AA1002103	9.64	4.31	6.49	12.68	13.50	19.90	7.83	9.63	9.63	*	+	
	Y79AA1002115	6.16	3.44	3.46	8.76	8.88	8.21	5.06	7.31	7.31	*	+	
	Y79AA1002121	4.13	1.90	2.75	5.52	3.99	4.66	2.99	2.94	2.94			
	Y79AA1002125	12.29	7.02	6.63	8.98	11.00	7.52	5.97	9.22	9.22			
	Y79AA1002129	4.01	2.55	2.79	4.98	5.25	5.00	4.03	4.07	4.07	*	+	
30	Y79AA1002131	3.98	1.83	2.10	2.08	2.08	3.32	2.24	4.89	4.89			
	Y79AA1002139	1.73	1.39	1.53	2.67	1.39	3.06	1.75	4.33	4.33			
	Y79AA1002144	13.61	9.16	11.69	45.27	42.86	41.51	20.24	31.90	31.90	**	+	*
	Y79AA1002177	11.17	7.99	8.29	8.46	8.96	11.14	8.89	10.57	10.57			
	Y79AA1002183	20.7	16.65	16.79	14.07	13.54	11.10	9.93	9.44	9.44	*	-	**
35	Y79AA1002202	16.44	8.10	6.76	14	14.11	9.08	7.13	7.42	7.42			
	Y79AA1002204	6.31	4.49	4.52	4.3	4.77	3.13	5.10	6.00	6			
	Y79AA1002206	3.17	2.15	1.77	3.09	3.03	2.45	3.04	3.50	3.5			
	Y79AA1002208	5.15	2.57	2.96	5.99	4.60	5.97	4.50	4.63	4.63			
	Y79AA1002209	3.58	4.01	5.76	4.15	3.13	3.39	4.99	7.55	7.55			
	Y79AA1002210	3.18	1.43	2.37	3.02	2.02	1.71	2.10	2.41	2.41			
40	Y79AA1002211	4.91	3.46	4.17	4.11	5.81	4.91	5.34	5.38	5.38		*	+
	Y79AA1002213	3.71	2.49	1.89	7.09	8.26	4.18	2.61	4.10	4.1	*	+	
	Y79AA1002215	12.98	6.72	6.55	11.46	10.70	7.31	10.62	11.29	11.29			
	Y79AA1002220	3.6	0.24	1.50	2.1	2.24	1.13	3.21	3.17	3.17			
	Y79AA1002226	15.84	9.35	12.55	20.91	22.33	23.57	11.78	20.18	20.18	**	+	
45	Y79AA1002229	6.49	3.85	3.45	4.63	4.19	3.44	5.38	5.16	5.16			
	Y79AA1002234	3.86	2.44	4.84	4.04	4.91	5.32	5.97	5.64	5.64		*	+
	Y79AA1002235	1.93	0.75	1.35	1.7	1.20	2.27	2.65	2.05	2.05			
	Y79AA1002246	2.63	2.09	2.74	2.5	3.97	3.37	2.34	1.90	1.9			
	Y79AA1002258	3.31	3.27	3.40	4.93	5.68	4.75	4.02	4.20	4.2	**	+	**
	Y79AA1002279	4.56	2.57	2.26	6.13	5.09	3.71	4.81	5.29	5.29			
50	Y79AA1002292	6.26	3.04	2.73	4.57	5.32	3.62	2.90	5.67	5.67			
	Y79AA1002298	1.82	0.51	1.99	1.65	1.57	1.30	1.17	0.87	0.87			
	Y79AA1002307	5.23	1.97	1.83	2.94	3.94	2.54	2.69	3.59	3.59			
	Y79AA1002309	1.73	1.34	1.76	1.52	3.43	2.98	1.67	1.76	1.76			
	Y79AA1002311	4.03	2.76	3.87	5	3.49	3.66	2.61	6.69	6.69			
55	Y79AA1002334	2.47	4.14	2.46	2.65	3.80	4.63	2.18	3.21	3.21			
	Y79AA1002351	3.38	3.58	4.03	5.8	3.67	5.58	3.56	6.63	6.63			
	Y79AA1002355	7.23	3.03	2.25	63.68	74.46	52.07	46.06	44.44	44.44	**	+	**

Table 347

5	Y79AA1001548	10.61	7.08	4.15	16.42	14.68	15.82	9.30	9.38	9.38	*	+		
	Y79AA1001555	7.52	5.37	3.80	6.53	5.95	5.70	7.04	7.00	7				
	Y79AA1001562	13.12	10.40	12.01	18.73	17.97	15.42	12.97	18.83	18.83	*	+		
	Y79AA1001581	2.59	2.12	1.33	2.27	2.33	1.95	1.31	2.40	2.4				
	Y79AA1001585	1.89	1.52	2.52	3.13	3.14	3.51	2.68	3.89	3.89	*	+	*	+
	Y79AA1001592	8.75	5.76	6.22	9.06	9.03	12.16	6.95	10.71	10.71				
10	Y79AA1001594	2.44	2.99	2.99	4.89	6.76	6.84	2.08	3.52	3.52	**	+		
	Y79AA1001603	41.01	29.22	27.39	35.33	47.15	41.79	19.68	22.24	22.24				
	Y79AA1001613	11.06	8.37	6.50	10.25	10.82	7.55	6.69	6.52	6.52				
	Y79AA1001630	0.95	0.54	0.85	1.19	0.72	0.95	1.19	0.88	0.88				
	Y79AA1001647	6.2	2.96	3.68	2.82	5.76	5.40	3.17	4.07	4.07				
15	Y79AA1001664	13.85	6.76	7.31	10.57	12.90	8.91	7.51	7.68	7.68				
	Y79AA1001665	3.6	3.81	4.37	4.15	4.52	5.51	3.17	4.23	4.23				
	Y79AA1001679	14	9.57	9.87	11.81	14.25	13.41	7.94	7.63	7.63				
	Y79AA1001692	3.06	2.79	3.66	3.62	3.64	6.60	2.78	2.76	2.76				
	Y79AA1001696	0.47	0.94	0.29	1.8	1.18	2.00	1.48	1.81	1.81	*	+	**	+
20	Y79AA1001705	5.59	4.16	3.52	5.12	5.14	5.00	3.05	4.02	4.02				
	Y79AA1001711	17.19	10.51	9.53	37.34	40.06	24.12	26.85	27.39	27.39	*	+	**	+
	Y79AA1001717	1.38	0.95	0.69	2.28	1.17	1.95	0.86	2.01	2.01				
	Y79AA1001719	3.1	2.90	1.65	4.96	4.48	2.69	2.06	2.48	2.48				
	Y79AA1001727	5.47	4.87	4.29	8.17	8.05	7.12	4.94	6.45	6.45	**	+		
25	Y79AA1001750	20.76	27.54	23.83	38.95	38.37	32.83	22.83	25.62	25.62	*	+		
	Y79AA1001760	6.22	6.83	3.78	10.14	8.09	8.51	8.09	4.11	4.11	*	+		
	Y79AA1001777	4.19	4.98	4.30	10.69	9.61	8.63	5.89	5.49	5.49	**	+	*	+
	Y79AA1001781	1.41	(0.02)	0.49	0.49	0.41	1.88	0.28	0.56	0.56				
	Y79AA1001787	6.73	4.26	4.09	6.64	5.23	7.45	4.25	5.24	5.24				
	Y79AA1001793	7.3	4.12	4.31	5.83	5.04	3.68	5.12	4.48	4.48				
30	Y79AA1001795	3	0.80	2.09	2.69	3.85	3.29	1.73	3.18	3.18				
	Y79AA1001799	5.26	2.91	2.67	5.21	5.65	6.10	3.13	5.77	5.77				
	Y79AA1001800	4.16	2.57	3.82	5.16	2.55	3.90	3.53	6.79	6.79				
	Y79AA1001801	6.56	3.89	3.46	8.87	3.49	7.02	3.18	4.68	4.68				
	Y79AA1001803	6.72	4.12	3.95	5.51	7.22	5.68	5.48	5.55	5.55				
35	Y79AA1001805	22.35	9.91	10.35	15.2	27.86	21.20	9.25	13.14	13.14				
	Y79AA1001807	6.96	2.99	4.40	6.3	4.51	3.72	4.95	5.25	5.25				
	Y79AA1001827	8.38	3.69	5.67	7.55	7.81	11.23	9.11	12.46	12.46		*	+	
	Y79AA1001846	4.45	2.15	3.75	6.2	4.92	5.41	3.96	7.82	7.82				
	Y79AA1001848	2.85	1.48	2.40	3.01	2.43	2.61	2.57	2.46	2.46				
40	Y79AA1001853	13.89	10.72	11.89	14.4	8.43	13.46	12.95	13.31	13.31				
	Y79AA1001863	15.14	7.58	9.41	15.02	11.89	14.02	8.02	12.33	12.33				
	Y79AA1001866	9.57	4.75	5.85	11.97	24.49	9.54	5.28	9.21	9.21				
	Y79AA1001874	1.66	0.73	0.26	0.48	1.10	0.61	0.67	0.63	0.63				
	Y79AA1001875	9	6.56	7.74	8.02	10.17	8.54	9.22	11.36	11.36		*	+	
45	Y79AA1001907	117.42	47.16	76.24	98.59	98.47	94.40	33.03	51.77	51.77				
	Y79AA1001908	2.02	0.84	1.62	1.52	1.88	1.08	1.03	1.18	1.18				
	Y79AA1001923	4.54	1.74	1.64	1.87	1.96	1.62	3.56	1.90	1.9				
	Y79AA1001927	7.1	4.39	6.61	6.81	4.65	6.65	7.02	7.63	7.63				
	Y79AA1001930	11.14	5.72	8.19	8.38	8.79	9.40	6.07	5.53	5.53				
	Y79AA1001932	4.55	2.74	2.35	3.75	4.07	2.57	2.61	2.55	2.55				
50	Y79AA1001933	5.44	2.77	4.71	4.94	5.23	2.92	4.54	4.08	4.08				
	Y79AA1001942	5.27	2.57	3.54	3.47	2.89	2.27	3.23	4.03	4.03				
	Y79AA1001963	16.6	6.83	11.89	12.17	15.10	11.74	8.04	11.18	11.18				
	Y79AA1001968	19.06	9.81	14.73	19.14	20.00	14.84	13.89	19.22	19.22				
	Y79AA1001983	8.12	2.93	5.67	4.13	3.77	4.79	3.53	4.78	4.78				
55	Y79AA1002000	8.2	3.32	3.60	7.79	5.70	5.11	5.48	4.05	4.05				
	Y79AA1002004	31.21	14.32	20.67	18.62	19.17	22.54	12.67	21.19	21.19				
	Y79AA1002008	6.53	5.69	5.73	9.64	7.46	8.70	4.99	6.08	6.08	*	+		

Table 346

	Y79AA1001077	11.3	7.81	9.27	10.02	10.61	11.75	11.20	11.01	11.01			
5	Y79AA1001078	2.85	2.15	2.01	4.62	7.48	2.90	4.22	3.26	3.26		*	+
	Y79AA1001081	16.61	9.85	12.79	10	10.38	11.30	5.81	7.08	7.08		*	-
	Y79AA1001088	26.22	15.63	20.41	21.72	24.28	26.25	25.14	31.31	31.31			
	Y79AA1001089	11.17	5.53	8.30	9.49	6.56	8.41	9.43	10.79	10.79			
	Y79AA1001090	4.51	2.54	4.20	6.81	5.20	6.61	4.39	5.95	5.95	*	+	
10	Y79AA1001105	27.01	7.71	19.38	6.68	4.28	6.75	6.37	6.27	6.27			
	Y79AA1001142	8.95	5.63	7.03	5.98	7.11	5.88	10.76	13.80	13.8		*	+
	Y79AA1001145	11.65	9.12	8.63	15.01	11.35	17.02	8.48	10.99	10.99			
	Y79AA1001162	4.06	1.39	1.51	5.09	3.87	3.44	4.59	3.13	3.13			
	Y79AA1001167	7.25	3.07	2.49	5.01	3.56	4.46	3.63	5.24	5.24			
15	Y79AA1001176	4.11	2.23	2.70	4.09	2.43	5.22	2.25	2.60	2.6			
	Y79AA1001177	4.68	4.25	4.38	3.59	3.61	5.91	4.61	3.71	3.71			
	Y79AA1001179	21.68	16.62	20.48	11.99	9.19	16.21	8.81	11.14	11.14	*	-	**
	Y79AA1001185	5.31	2.79	3.61	5.39	3.59	5.46	3.84	4.29	4.29			
	Y79AA1001201	28.52	17.14	23.93	16.35	22.62	37.53	18.59	26.16	26.16			
20	Y79AA1001205	10.97	3.75	3.90	5.2	4.84	4.63	3.49	3.72	3.72			
	Y79AA1001211	11.99	5.80	6.48	8.33	12.82	9.17	4.23	4.74	4.74			
	Y79AA1001212	7.31	3.41	4.24	5.88	4.00	4.88	4.13	6.49	6.49			
	Y79AA1001216	55.35	32.24	33.00	52.32	49.82	57.61	27.61	40.72	40.72			
	Y79AA1001228	9.47	5.39	6.44	9.83	8.83	13.70	14.26	14.88	14.88		**	+
	Y79AA1001233	7.94	5.13	5.27	5.47	5.22	5.58	6.11	7.96	7.96			
25	Y79AA1001236	9.41	4.91	6.23	8.19	6.64	8.01	4.19	7.99	7.99			
	Y79AA1001239	17.51	11.16	12.48	23.85	15.23	20.67	15.26	22.26	22.26			
	Y79AA1001240	6.74	4.58	4.53	7.09	6.25	7.67	6.30	7.17	7.17			
	Y79AA1001255	11.62	4.94	6.87	6.84	9.34	6.89	3.77	5.35	5.35			
	Y79AA1001264	8.92	4.36	4.37	5.15	4.83	5.09	6.25	11.76	11.76			
30	Y79AA1001272	16.07	9.52	9.48	17.58	13.84	18.59	12.50	13.21	13.21			
	Y79AA1001281	2.39	1.46	1.20	2.86	1.50	1.94	1.67	2.71	2.71			
	Y79AA1001299	15.84	12.69	13.71	17.01	14.77	25.21	17.79	21.80	21.8		*	+
	Y79AA1001312	7.69	3.18	3.48	9.46	10.75	7.56	6.31	5.09	5.09			
	Y79AA1001319	9.18	6.58	8.51	11.43	8.41	10.88	8.28	9.95	9.95			
35	Y79AA1001323	5.8	3.74	3.41	4.67	5.59	4.56	4.04	5.77	5.77			
	Y79AA1001328	9.21	5.33	4.01	6.44	6.42	8.24	6.73	9.42	9.42			
	Y79AA1001343	862.89	462.45	576.89	529.68	551.94	571.68	1081.07	1529.21	1529		*	+
	Y79AA1001351	1.98	0.57	1.69	0.7	1.23	1.95	1.38	2.51	2.51			
	Y79AA1001364	13.67	8.79	10.09	17.42	16.54	19.67	6.03	14.83	14.83	*	+	
40	Y79AA1001367	6.28	4.16	4.34	5.94	4.67	6.56	4.76	4.90	4.9			
	Y79AA1001384	1.87	1.73	1.53	1.86	1.16	2.08	1.66	1.46	1.46			
	Y79AA1001391	3.6	2.56	1.82	3.57	3.95	4.39	3.23	2.67	2.67			
	Y79AA1001394	7.58	3.85	2.91	6.13	4.47	4.34	2.98	3.74	3.74			
	Y79AA1001402	14.12	9.28	8.02	15.91	14.24	20.22	15.90	16.49	16.49		*	+
45	Y79AA1001410	6.61	3.47	3.47	4.7	4.77	5.26	4.23	5.49	5.49			
	Y79AA1001414	4.82	2.47	3.52	4.85	3.10	4.46	3.68	4.21	4.21			
	Y79AA1001426	6.98	4.46	5.28	4.95	5.72	4.24	5.87	6.84	6.84			
	Y79AA1001427	3.95	3.35	3.13	5.95	6.19	3.76	4.11	6.23	6.23			
	Y79AA1001430	3.36	4.23	3.56	4.36	4.28	5.52	6.35	7.62	7.62		**	+
	Y79AA1001439	4.05	2.77	2.23	5.27	3.53	5.80	5.59	7.03	7.03		**	+
50	Y79AA1001485	1.52	0.56	1.47	1.8	1.02	1.80	1.03	1.44	1.44			
	Y79AA1001493	1.38	0.86	0.94	2.07	2.04	2.30	1.06	3.12	3.12	**	+	
	Y79AA1001511	7.88	6.25	4.30	5.78	8.49	6.85	6.79	10.70	10.7			
	Y79AA1001523	10.75	7.00	5.10	7.67	3.96	7.11	7.19	5.22	5.22			
	Y79AA1001530	6.54	3.62	3.97	4.77	5.40	7.25	5.26	7.92	7.92			
55	Y79AA1001532	4.4	3.73	3.24	7.17	5.68	6.14	4.71	4.96	4.96	*	+	*
	Y79AA1001533	5.01	4.00	3.23	3.96	6.97	7.08	2.96	4.09	4.09			
	Y79AA1001541	12.19	9.13	10.66	12.21	12.80	16.01	5.59	5.41	5.41		**	-

Table 345

5	Y79AA1000724	28.17	13.18	13.80	13.88	13.98	11.98	3.06	4.28	4.28		*	-
	Y79AA1000726	8.11	5.46	4.24	6.09	4.77	4.52	5.43	7.82	7.82			
	Y79AA1000734	3.88	2.62	-2.34	5.17	3.55	4.31	2.92	6.05	6.05			
	Y79AA1000748	3.95	1.81	1.83	2.64	2.02	2.92	1.57	2.24	2.24			
	Y79AA1000750	10.39	6.10	4.86	9.81	8.59	9.78	5.43	7.43	7.43			
	Y79AA1000752	2.87	0.53	1.08	2.54	2.81	2.11	1.32	1.59	1.59			
10	Y79AA1000774	5.72	4.59	2.86	2.14	2.79	5.77	3.53	3.76	3.76			
	Y79AA1000776	4.35	4.36	2.86	3.71	4.12	5.01	3.48	3.30	3.3			
	Y79AA1000777	11.76	6.21	5.54	8.56	11.90	10.17	6.16	6.66	6.66			
	Y79AA1000778	13.22	6.87	8.41	14.77	13.90	13.40	7.19	13.72	13.72			
	Y79AA1000782	7.86	4.93	5.51	5.52	4.90	5.05	5.46	7.23	7.23			
15	Y79AA1000784	12.43	9.12	11.59	13	14.52	14.46	11.05	11.31	11.31			
	Y79AA1000794	4.35	2.95	2.89	4.43	4.95	3.90	3.24	3.10	3.1			
	Y79AA1000800	2.57	2.36	2.08	3	3.32	3.30	2.93	3.69	3.69	**	+	+
	Y79AA1000802	1.85	1.48	1.65	1	0.76	1.64	0.34	1.23	1.23			
	Y79AA1000805	4.24	3.55	2.28	3.22	3.19	3.89	2.71	4.15	4.15			
	Y79AA1000814	14.61	9.83	7.28	9.51	9.83	6.77	3.86	4.30	4.3		*	-
20	Y79AA1000823	12.6	9.53	9.56	15.44	14.21	12.23	9.08	15.12	15.12			
	Y79AA1000824	4.44	3.44	2.16	2.49	3.58	2.72	2.72	3.74	3.74			
	Y79AA1000827	3.1	1.46	1.84	2.99	1.29	1.77	1.89	2.61	2.61			
	Y79AA1000831	5.49	4.85	5.37	3.74	4.89	3.85	3.76	5.38	5.38			
	Y79AA1000833	40.22	31.45	37.17	40.96	46.51	50.53	34.20	40.04	40.04			
25	Y79AA1000850	2.09	2.81	2.57	4.27	3.76	4.02	3.33	2.26	2.26	**	+	
	Y79AA1000856	6.74	5.50	6.27	7.85	6.17	10.60	4.73	5.48	5.48			
	Y79AA1000862	12.52	7.78	4.39	13.89	9.86	8.13	7.63	7.94	7.94			
	Y79AA1000876	8.46	4.16	4.01	6.87	6.89	6.26	3.75	5.07	5.07			
	Y79AA1000888	1.47	1.34	1.40	1.56	1.46	1.29	1.98	1.99	1.99		**	+
30	Y79AA1000902	16.38	10.81	14.11	11.4	9.46	11.97	5.88	7.23	7.23		*	-
	Y79AA1000935	16.25	11.98	13.09	25.37	21.17	25.92	23.44	29.28	29.28	**	+	**
	Y79AA1000959	3.1	2.66	3.26	3.18	3.69	2.84	2.68	4.50	4.5			
	Y79AA1000962	1.8	2.34	1.77	4.45	3.80	4.94	2.33	2.34	2.34	**	+	
	Y79AA1000963	43.49	20.23	23.14	40.9	40.35	45.98	17.97	19.24	19.24			
35	Y79AA1000966	8	6.62	3.05	7.53	7.98	4.56	6.48	5.59	5.59			
	Y79AA1000967	11.14	8.37	5.21	15.29	15.02	10.80	8.86	10.67	10.67			
	Y79AA1000968	11.05	6.63	3.78	6.32	9.03	6.81	4.66	7.08	7.08			
	Y79AA1000969	4.13	3.63	3.19	4.09	3.12	3.96	2.88	4.11	4.11			
	Y79AA1000976	2.07	1.66	1.63	2.46	2.43	2.76	2.15	3.14	3.14	*	+	*
	Y79AA1000978	3.15	2.68	2.59	3.19	2.43	2.99	1.56	2.57	2.57			
40	Y79AA1000985	4.53	6.21	3.11	9.92	6.66	7.93	4.84	4.19	4.19			
	Y79AA1000989	27.14	18.46	21.17	22.61	22.40	25.64	17.86	17.83	17.83			
	Y79AA1000991	14.41	7.65	8.70	14.5	16.91	8.11	10.68	10.04	10.04			
	Y79AA1001013	35.7	19.64	14.11	24.63	29.38	32.01	18.46	27.65	27.65			
	Y79AA1001014	8.41	5.13	3.58	6.96	7.27	8.35	6.51	8.47	8.47			
45	Y79AA1001019	6.41	3.32	4.05	4.98	4.88	5.75	4.58	5.04	5.04			
	Y79AA1001020	13.26	4.81	6.74	9.29	9.05	11.19	6.66	10.83	10.83			
	Y79AA1001023	3.99	2.27	3.29	3.71	4.41	3.42	4.24	3.90	3.9			
	Y79AA1001030	4.36	2.82	3.64	7.73	6.53	9.26	7.69	8.68	8.68	**	+	**
	Y79AA1001035	-0.01	7.25	7.50	9.11	6.84	10.21	7.88	15.95	15.95			
50	Y79AA1001041	8.33	4.39	3.51	5.69	4.65	4.21	2.70	5.79	5.79			
	Y79AA1001043	18.02	12.74	10.03	8.74	8.44	9.66	9.73	11.39	11.39			
	Y79AA1001048	5.98	4.37	5.02	5.57	4.35	5.46	5.24	5.86	5.86			
	Y79AA1001056	2.8	1.67	2.69	4.83	3.64	3.93	2.91	3.52	3.52	*	+	
	Y79AA1001061	4.66	2.07	2.99	8.42	5.16	8.18	3.38	5.08	5.08	*	+	
	Y79AA1001062	4.59	1.72	3.28	8.74	7.23	9.33	3.95	5.55	5.55	**	+	
55	Y79AA1001068	7.33	4.55	5.57	10.85	9.20	12.48	5.68	6.64	6.64	*	+	
	Y79AA1001073	12.4	6.75	7.01	7.75	5.93	9.79	5.80	7.72	7.72			

Table 344

5	Y79AA1000280	11.25	5.37	6.77	11.8	13.92	12.66	5.46	9.54	9.54			
	Y79AA1000285	4.46	1.52	2.70	3.31	1.78	2.60	2.43	3.53	3.53			
	Y79AA1000295	3.61	2.65	~9.31	10.15	10.34	10.77	4.41	5.66	5.66	**	+	* +
	Y79AA1000307	12.46	9.65	13.13	11.87	8.54	13.75	5.29	6.68	6.68		**	-
	Y79AA1000313	15.46	6.94	8.62	10.28	12.44	14.87	10.41	13.90	13.9			
	Y79AA1000314	14.81	9.18	10.30	22.74	18.92	27.80	24.11	31.46	31.46	*	+	** +
10	Y79AA1000328	3.09	1.87	2.24	2.09	2.55	2.73	1.78	2.96	2.96			
	Y79AA1000334	7.09	3.70	2.56	5.55	4.48	4.69	3.41	4.25	4.25			
	Y79AA1000342	35.87	15.66	15.62	22.36	17.70	23.91	21.00	29.07	29.07			
	Y79AA1000346	17.41	15.57	12.74	9.41	9.10	10.71	4.23	5.49	5.49	*	-	** -
	Y79AA1000347	23.11	14.24	15.07	23.5	39.38	38.47	19.81	25.73	25.73	*	+	
15	Y79AA1000349	19.76	10.53	12.68	20.31	16.01	21.05	12.82	17.27	17.27			
	Y79AA1000355	4.87	2.42	3.06	7.26	6.44	8.31	4.76	6.17	6.17	*	+	
	Y79AA1000368	6.76	2.87	3.15	4.62	3.69	5.41	4.31	4.40	4.4			
	Y79AA1000388	25.23	15.44	16.71	26.79	21.25	29.10	12.60	17.85	17.85			
	Y79AA1000392	14.91	8.34	19.71	13.34	7.02	19.13	9.61	11.82	11.82			
20	Y79AA1000405	24.03	14.82	7.15	15.39	22.71	12.76	14.12	17.35	17.35			
	Y79AA1000410	24.25	16.23	12.97	37.19	36.14	36.35	20.62	22.06	22.06	**	+	
	Y79AA1000420	1.83	1.06	1.88	2.33	1.74	3.81	1.85	2.84	2.84			
	Y79AA1000423	7.25	4.11	5.48	9.75	7.86	8.44	5.00	5.45	5.45	*	+	
	Y79AA1000426	5.29	3.84	5.55	4.45	2.88	4.33	3.32	3.94	3.94			
25	Y79AA1000432	3.27	2.71	3.28	1.62	1.68	2.55	1.63	2.22	2.22	*	-	* -
	Y79AA1000453	141.24	53.68	107.37	81.71	59.38	81.50	30.05	43.77	43.77			
	Y79AA1000465	3.59	1.59	2.02	2.43	1.32	2.55	1.95	3.10	3.1			
	Y79AA1000469	14.01	11.65	7.90	12.08	10.53	7.10	8.31	7.33	7.33			
	Y79AA1000480	4.69	1.58	1.60	4.05	2.82	2.60	2.60	2.44	2.44			
	Y79AA1000502	12.81	5.39	8.31	9.83	13.49	9.32	5.96	11.12	11.12			
30	Y79AA1000521	6.28	4.42	6.32	6.26	4.77	4.40	6.38	6.79	6.79			
	Y79AA1000534	17.26	8.63	8.69	10.74	7.23	7.43	4.39	5.56	5.56			
	Y79AA1000538	6.63	3.28	4.52	10.32	7.26	8.06	5.36	6.47	6.47	*	+	
	Y79AA1000539	19.25	8.27	12.78	24.31	26.47	21.68	9.27	11.72	11.72	*	+	
	Y79AA1000540	11.13	5.92	6.15	9.13	9.09	8.44	6.65	9.21	9.21			
35	Y79AA1000560	173.06	134.34	94.53	202.66	161.69	169.55	95.78	139.04	139			
	Y79AA1000574	2.89	2.45	2.28	4.12	2.97	2.60	1.96	2.63	2.63			
	Y79AA1000584	3.2	1.68	1.63	1.75	2.10	2.56	2.05	2.41	2.41			
	Y79AA1000589	8.66	5.80	5.36	6.79	3.71	6.73	6.49	7.62	7.62			
	Y79AA1000598	5.98	2.97	4.18	3.57	3.29	6.10	4.35	4.63	4.63			
40	Y79AA1000600	6.57	3.44	3.89	3.3	2.25	3.48	2.55	2.77	2.77			
	Y79AA1000609	6.92	3.42	2.75	2.76	4.04	6.09	4.13	5.52	5.52			
	Y79AA1000618	58.41	30.55	40.08	29.92	37.02	38.12	11.43	14.49	14.49		*	-
	Y79AA1000627	6.08	3.22	3.45	5.69	5.50	4.18	4.40	3.93	3.93			
	Y79AA1000636	38.19	23.55	23.75	16.84	22.87	15.14	9.44	11.05	11.05		*	-
45	Y79AA1000649	8.69	4.34	4.67	4.61	4.61	4.01	3.93	8.79	8.79			
	Y79AA1000656	5.76	3.08	3.22	5.58	4.90	5.16	3.04	4.23	4.23			
	Y79AA1000673	5.03	2.72	1.36	3.23	1.94	2.41	3.39	4.06	4.06			
	Y79AA1000674	10.61	7.11	11.17	10.18	8.67	10.62	6.76	10.00	10			
	Y79AA1000678	7.25	4.89	6.06	10.19	6.81	7.33	5.06	5.92	5.92			
	Y79AA1000682	24.87	16.17	18.30	22.46	26.14	12.58	16.87	20.51	20.51			
50	Y79AA1000683	15.32	7.96	8.21	6.64	7.38	6.63	4.88	6.13	6.13			
	Y79AA1000697	54.8	30.85	37.16	42.84	41.24	37.90	36.31	42.61	42.61			
	Y79AA1000700	9.78	3.97	5.64	3.6	3.66	3.51	4.90	7.29	7.29			
	Y79AA1000702	17.82	9.90	9.05	9.33	10.05	8.49	5.94	9.28	9.28			
	Y79AA1000704	2.05	0.88	1.35	0.9	1.80	0.91	1.41	1.66	1.66			
	Y79AA1000705	2.45	1.63	1.24	3.99	2.73	3.50	2.26	2.26	2.26	*	+	
55	Y79AA1000717	11.47	6.51	7.68	14.26	8.53	10.93	5.99	11.28	11.28			
	Y79AA1000722	6.59	5.15	4.02	3.83	3.18	4.48	1.26	1.65	1.65		**	-

Table 343

5	VESEN2000272	6.36	2.52	3.61	13.68	15.50	9.23	6.37	6.11	6.11	*	+		
	VESEN2000299	5.8	3.32	3.03	6.33	5.54	5.31	4.11	3.82	3.82				
	VESEN2000323	3.64	2.70	3.46	7.25	6.60	6.83	4.13	6.99	6.99	**	+	*	+
	VESEN2000327	16.91	9.24	9.32	14.89	11.98	16.05	16.51	12.53	12.53				
	VESEN2000328	3.41	1.69	2.05	2.7	1.99	2.52	3.68	4.21	4.21			*	+
10	VESEN2000330	9.06	4.94	3.98	4	3.94	4.40	7.56	5.58	5.58				
	VESEN2000336	3.29	2.35	2.63	3.19	2.56	2.84	2.06	2.38	2.38				
	VESEN2000354	8.7	4.46	4.22	7.46	6.89	5.83	5.63	5.02	5.02				
	VESEN2000378	3.42	2.15	2.25	4.13	2.42	1.91	1.92	2.61	2.61				
	VESEN2000379	11.63	7.79	4.82	10.74	10.07	12.49	7.29	10.70	10.7				
	VESEN2000397	3.37	1.29	1.36	2.39	2.24	1.99	1.18	3.19	3.19				
15	VESEN2000416	3.83	2.34	1.55	2.15	2.33	2.91	2.47	2.28	2.28				
	VESEN2000420	2.88	0.98	1.36	1.52	0.23	0.52	0.64	1.63	1.63				
	VESEN2000430	2.62	1.65	1.71	1.89	2.49	1.83	0.78	2.97	2.97				
	VESEN2000448	2.86	2.67	1.17	1	2.01	2.37	2.33	2.73	2.73				
	VESEN2000449	8.25	5.92	4.67	9.14	8.56	10.89	5.16	6.55	6.55				
20	VESEN2000456	5.37	3.06	1.86	3.12	2.41	3.57	2.05	2.65	2.65				
	VESEN2000562	7.78	4.41	5.30	5.84	5.51	4.92	4.30	6.44	6.44				
	VESEN2000573	0.6	0.35	0.41	0.67	0.40	0.67	1.28	2.60	2.6			*	+
	VESEN2000604	5.64	1.48	1.85	3.25	2.37	2.19	2.91	4.05	4.05				
	VESEN2000614	25.21	13.24	16.03	20.97	19.46	20.96	23.97	21.61	21.61				
25	VESEN2000638	1.7	1.28	1.62	1.56	1.85	1.20	2.41	1.35	1.35				
	VESEN2000641	1.73	2.11	1.08	1.79	1.66	1.77	1.14	1.95	1.95				
	VESEN2000645	3.09	2.77	2.30	2.12	2.14	1.71	1.70	3.15	3.15				
	Y79AA1000013	10.79	7.40	5.68	11.91	9.74	8.63	7.82	6.74	6.74				
	Y79AA1000030	13.95	8.47	8.24	10.96	9.10	13.62	9.47	12.29	12.29				
30	Y79AA1000033	16.96	12.16	9.55	7.65	10.20	8.44	7.18	10.76	10.76				
	Y79AA1000037	2.11	1.49	0.71	2.23	2.21	3.27	2.75	2.51	2.51			*	+
	Y79AA1000041	2.2	2.48	1.77	2.69	2.36	2.74	2.02	3.82	3.82				
	Y79AA1000059	7.6	6.90	6.65	10.99	11.69	12.90	4.30	7.70	7.7	**	+		
	Y79AA1000065	22.39	17.36	15.96	24.43	21.67	25.09	14.43	16.06	16.06				
	Y79AA1000081	42.69	41.35	51.24	111.38	113.45	103.25	45.62	16.30	16.3	**	+		
35	Y79AA1000127	22.29	16.01	11.79	12.57	10.65	7.07	3.98	5.58	5.58			*	-
	Y79AA1000130	6.17	3.27	2.80	10.01	8.60	9.63	4.89	5.13	5.13	**	+		
	Y79AA1000131	448.19	235.19	299.39	399.75	486.28	438.12	277.73	304.61	304.6				
	Y79AA1000134	8.96	7.49	5.25	6.6	6.53	6.62	9.23	10.69	10.69				
	Y79AA1000143	9.99	4.29	8.06	7.58	8.06	8.95	6.96	8.30	8.3				
40	Y79AA1000144	8.55	7.18	6.04	6.31	5.55	6.00	4.05	4.40	4.4			*	-
	Y79AA1000150	18.22	14.18	15.26	14.89	15.33	21.06	9.92	9.91	9.91			**	-
	Y79AA1000153	201.67	139.66	172.85	190.71	189.25	179.30	103.81	119.17	119.2			*	-
	Y79AA1000166	6.51	3.61	2.42	6.7	8.84	4.48	3.56	4.21	4.21				
	Y79AA1000179	15.16	9.65	7.92	10.53	9.30	7.94	4.29	5.64	5.64				
	Y79AA1000181	10.66	5.30	5.63	7.26	8.22	5.85	3.94	5.98	5.98				
45	Y79AA1000202	18.5	15.06	12.86	18.25	18.73	23.11	15.84	25.98	25.98				
	Y79AA1000207	5.87	4.02	4.27	14.67	14.22	14.10	7.48	5.85	5.85	**	+		
	Y79AA1000214	29.22	23.27	20.29	36.32	37.06	45.40	22.86	25.86	25.86	*	+		
	Y79AA1000222	12.84	9.84	10.93	9.21	6.89	9.06	5.29	5.66	5.66			**	-
	Y79AA1000226	5.63	6.09	5.68	7.41	7.20	8.09	8.84	8.79	8.79	**	+	**	+
50	Y79AA1000227	17.27	10.05	8.43	12.69	17.80	12.32	9.20	10.19	10.19				
	Y79AA1000230	6.42	4.02	2.20	3.72	4.88	2.48	3.03	2.90	2.9				
	Y79AA1000231	34.72	21.74	21.36	20.87	19.10	17.13	9.21	15.10	15.1				
	Y79AA1000239	15.79	9.79	7.30	10.27	13.40	11.55	12.77	13.82	13.82				
	Y79AA1000258	4.05	3.20	3.26	4.22	5.80	4.84	3.99	4.25	4.25				
55	Y79AA1000268	7.27	4.70	4.79	10.11	6.83	6.96	5.20	6.24	6.24				
	Y79AA1000269	3.42	2.81	2.55	4.54	6.08	5.88	5.38	5.60	5.6	**	+	**	+
	Y79AA1000270	3.64	4.17	2.51	5.74	6.14	5.66	3.62	4.41	4.41	**	+		

Table 342

5	THYRO1001828	5.58	5.56	4.00	9.32	9.83	9.03	4.86	6.29	6.29	**	+		
	THYRO1001854	20.22	7.97	7.27	24.83	26.41	23.02	14.19	14.50	14.5	*	+		
	THYRO1001895	4.5	1.82	1.66	2.69	3.40	3.20	2.51	2.17	2.17				
	THYRO1001907	6.37	2.87	2.77	7.43	8.35	6.14	3.08	4.67	4.67				
	TRACH1000006	1.82	2.19	1.60	2.9	3.42	2.53	2.58	3.05	3.05	*	+	*	+
	TRACH1000013	2.15	1.13	1.31	1.45	1.80	3.25	1.50	1.76	1.76				
10	TRACH1000074	3.42	3.57	4.39	5.62	7.83	7.88	4.19	10.27	10.27	*	+		
	TRACH1000095	2.45	2.91	2.44	3.1	3.04	4.04	2.50	2.45	2.45				
	TRACH1000102	7.43	5.84	4.56	10.07	11.80	13.53	5.10	8.65	8.65	*	+		
	TRACH1000108	3.15	1.08	0.60	4.55	2.50	3.75	3.10	1.49	1.49				
	TRACH1000126	6.59	4.83	4.15	6.73	6.75	6.24	2.66	4.52	4.52				
15	TRACH1000146	4.1	2.48	3.17	3.77	4.50	3.73	2.81	3.85	3.85				
	TRACH1000160	2.88	1.73	0.69	2.15	3.29	1.84	1.31	2.46	2.46				
	TRACH1000184	9.18	5.15	6.68	9.87	12.29	12.18	7.92	7.13	7.13	*	+		
	VESEN1000004	1.43	3.20	2.03	4.77	4.23	4.76	2.44	2.90	2.9	*	+		
	VESEN1000007	4.67	3.71	3.03	4.92	4.79	4.78	3.45	3.27	3.27				
20	VESEN1000013	3.8	4.40	3.49	6.08	5.11	8.39	4.08	5.78	5.78				
	VESEN1000028	10.32	4.13	4.71	9.23	9.35	9.07	7.29	12.27	12.27				
	VESEN1000059	7.75	3.60	4.26	7.63	6.94	7.73	4.60	5.95	5.95				
	VESEN1000100	14.3	7.29	8.52	11.77	17.29	16.55	10.06	12.85	12.85				
	VESEN1000107	8.09	2.86	4.55	5.28	4.93	5.96	5.50	6.28	6.28				
	VESEN1000117	4.56	2.53	3.13	3.83	3.21	3.98	3.40	4.83	4.83				
25	VESEN1000122	6	2.68	4.24	3.89	4.52	7.18	4.38	7.65	7.65				
	VESEN1000137	2.93	1.73	1.82	1.57	3.65	3.17	2.10	3.43	3.43				
	VESEN1000195	14.98	5.35	5.89	8.11	8.22	6.74	10.54	12.97	12.97				
	VESEN1000215	2.26	0.13	1.20	1.57	1.68	0.85	0.67	1.63	1.63				
	VESEN1000279	26.58	15.13	14.91	21.43	14.13	23.59	19.30	20.07	20.07				
30	VESEN1000363	15.34	8.73	10.79	17.48	16.61	12.88	9.72	13.31	13.31				
	VESEN1000388	9.91	6.40	6.52	7.89	4.01	10.40	6.86	10.14	10.14				
	VESEN1000394	12.12	6.72	8.23	12.56	8.96	9.43	5.04	9.23	9.23				
	VESEN1000410	10.78	2.59	2.39	6.85	3.24	4.07	5.06	8.94	8.94				
	VESEN1000411	6.18	3.27	4.03	5.74	3.11	6.71	4.21	5.31	5.31				
35	VESEN1000415	9.24	6.34	4.20	8.16	6.27	5.95	4.08	7.14	7.14				
	VESEN1000440	9.05	5.57	4.80	8.89	8.64	8.72	5.45	8.25	8.25				
	VESEN1000452	7.8	4.72	5.60	4.86	5.38	4.21	6.76	5.77	5.77				
	VESEN1000539	346.75	188.95	244.65	158.74	166.73	144.68	64.90	151.18	151.2				
	VESEN1000554	4.46	3.39	3.95	4.07	2.23	3.58	2.95	2.93	2.93		*	-	
40	VESEN1000557	6.06	4.00	4.41	6.38	3.08	5.06	6.10	7.77	7.77		*	+	
	VESEN1000575	7.82	4.18	4.70	6.03	4.15	4.58	5.87	6.64	6.64				
	VESEN1000585	9.14	4.16	5.29	6.86	6.14	7.55	4.21	6.93	6.93				
	VESEN1000592	1.51	0.34	0.06	1.48	0.81	0.75	1.11	0.98	0.98				
	VESEN1000658	9.42	5.35	3.63	6.6	8.13	5.18	7.65	9.88	9.88				
	VESEN1000669	30.52	16.02	17.70	27.74	22.51	23.12	18.76	27.04	27.04				
45	VESEN1000743	12.62	7.52	8.22	9.64	10.40	10.72	6.57	9.41	9.41				
	VESEN1000752	31.33	20.56	19.92	44.49	19.58	40.73	21.19	32.70	32.7				
	VESEN1000761	23.86	13.01	17.50	12.45	9.94	17.39	8.43	10.21	10.21				
	VESEN2000039	77.69	44.95	56.28	57.5	43.19	64.97	60.33	69.54	69.54				
	VESEN2000102	7.33	4.99	5.35	6.83	4.25	7.08	6.69	8.37	8.37				
50	VESEN2000164	5.18	3.46	3.31	9.13	9.21	6.82	3.36	3.89	3.89	*	+		
	VESEN2000175	1.73	0.97	0.12	1.01	1.92	1.13	0.88	1.17	1.17				
	VESEN2000186	19.39	12.37	11.60	17.79	16.80	18.96	15.99	20.01	20.01				
	VESEN2000199	28.49	19.51	19.01	18.68	21.47	33.21	23.58	23.01	23.01				
	VESEN2000200	6.32	1.63	3.02	5.06	3.00	3.70	3.04	4.39	4.39				
55	VESEN2000204	4.52	1.87	3.26	2.47	1.87	2.02	2.17	3.09	3.09				
	VESEN2000218	6.43	3.74	5.10	6.59	6.27	8.76	4.84	5.35	5.35				
	VESEN2000230	5.26	2.88	3.63	6.04	5.20	6.82	6.20	5.85	5.85		*	+	

Table 341

5	THYRO1001498	9.2	3.54	3.52	8.32	6.23	9.44	6.75	6.00	6				
	THYRO1001510	8.51	2.92	3.62	4.12	4.26	4.21	2.96	4.74	4.74				
	THYRO1001512	9.32	6.84	5.74	9.67	9.37	8.03	7.58	10.22	10.22				
	THYRO1001519	9.13	4.10	4.70	9.27	7.38	9.67	6.98	8.20	8.2				
	THYRO1001522	6.26	4.50	5.23	7.93	8.82	7.33	5.58	9.26	9.26*	+			
	THYRO1001523	3.53	2.10	1.99	6.46	5.54	6.24	4.04	4.29	4.29**	+	+		+
10	THYRO1001526	6.91	4.84	5.74	14.18	9.51	13.49	12.30	16.11	16.11*	+	**	+	
	THYRO1001529	2.41	1.14	1.41	2.28	1.58	4.28	2.24	2.20	2.2				
	THYRO1001534	3.65	2.24	1.50	4.38	3.58	6.43	2.88	4.21	4.21				
	THYRO1001537	18.2	10.50	9.67	21.59	21.38	19.81	8.19	10.14	10.14*	+			
	THYRO1001541	14.28	6.89	6.76	16.77	16.36	14.76	9.61	10.03	10.03				
15	THYRO1001545	3.56	2.76	2.72	3.42	3.96	4.48	3.96	4.30	4.3		*	+	
	THYRO1001559	3.99	2.04	2.13	4.24	3.76	7.51	3.56	3.91	3.91				
	THYRO1001563	11.96	7.39	6.70	7.96	5.68	9.41	7.19	8.07	8.07				
	THYRO1001570	4.68	4.47	3.76	4.09	3.00	4.87	4.64	6.87	6.87				
	THYRO1001573	8.02	5.52	6.21	6.26	3.61	8.28	6.11	6.00	6				
20	THYRO1001584	8.32	5.29	4.71	9.43	6.63	9.84	5.17	6.12	6.12				
	THYRO1001593	2.99	0.93	1.22	3.14	4.86	2.61	2.01	4.21	4.21				
	THYRO1001595	5.67	1.96	2.39	7.68	7.67	6.34	3.91	4.14	4.14*	+			
	THYRO1001596	5.89	2.66	3.80	3.78	3.65	3.11	2.98	3.57	3.57				
	THYRO1001602	7.81	2.64	3.23	7.32	8.69	7.89	4.74	7.00	7				
	THYRO1001605	5.26	2.56	2.24	5.13	5.05	4.87	3.48	3.41	3.41				
25	THYRO1001608	7.75	3.89	6.86	6.23	6.07	8.04	6.19	6.87	6.87				
	THYRO1001617	14.26	9.34	10.47	17.37	15.68	19.92	9.80	12.17	12.17*	+			
	THYRO1001634	4.95	3.06	3.93	4.4	3.84	4.30	4.75	4.39	4.39				
	THYRO1001637	10.18	6.14	4.65	17.45	14.38	17.46	8.06	9.17	9.17**	+			
	THYRO1001641	6.38	3.44	3.03	6.59	5.36	5.81	5.90	5.59	5.59				
30	THYRO1001656	4.52	2.95	2.83	3.81	4.14	7.31	4.33	5.14	5.14				
	THYRO1001658	4.29	2.01	1.79	2.18	2.89	2.10	2.16	2.58	2.58				
	THYRO1001661	3.1	1.45	1.64	1.96	2.33	1.46	4.01	2.50	2.5				
	THYRO1001671	5.77	2.59	2.20	4.22	4.26	4.64	3.03	5.39	5.39				
	THYRO1001672	6.81	4.51	5.53	5.21	5.27	6.87	6.28	6.63	6.63				
35	THYRO1001673	4	1.65	1.66	5.32	3.21	5.73	2.44	2.64	2.64				
	THYRO1001677	6.31	4.12	3.30	6.16	7.35	6.56	2.26	3.46	3.46				
	THYRO1001683	8.24	4.40	3.37	4.91	4.29	8.77	5.76	11.28	11.28				
	THYRO1001700	4.49	4.00	2.73	4.05	4.60	4.19	4.01	4.47	4.47				
	THYRO1001702	15.24	5.52	7.38	9.42	10.75	10.20	8.66	10.47	10.47				
	THYRO1001703	9.25	6.47	6.51	7.26	6.71	8.49	10.46	8.63	8.63				
40	THYRO1001706	4.3	2.92	3.16	5.43	6.68	7.52	2.62	4.78	4.78*	+			
	THYRO1001721	5.23	3.35	2.76	6.77	6.22	4.74	5.26	7.04	7.04		*	+	
	THYRO1001725	4.92	2.94	2.29	5.59	6.33	8.71	2.75	4.72	4.72*	+			
	THYRO1001730	24.29	13.18	13.43	14.02	17.03	13.70	21.66	22.76	22.76				
	THYRO1001738	9.75	4.90	4.82	9.04	5.85	7.43	4.92	7.98	7.98				
45	THYRO1001743	4	3.23	1.86	3.1	3.27	3.05	4.23	2.96	2.96				
	THYRO1001745	2.52	1.07	1.25	1.89	1.88	1.53	1.75	2.88	2.88				
	THYRO1001746	4.33	2.26	1.61	3.33	3.91	4.02	3.18	3.68	3.68				
	THYRO1001770	12.11	9.28	9.48	15.96	14.31	15.08	8.82	12.34	12.34**	+			
	THYRO1001772	5.17	2.74	2.39	6.3	7.93	7.50	3.31	3.90	3.9*	+			
50	THYRO1001778	15.04	12.42	11.52	14.35	19.15	15.23	13.02	16.18	16.18				
	THYRO1001793	14.69	6.79	7.43	12.47	11.95	9.29	6.11	8.93	8.93				
	THYRO1001796	11.97	7.13	5.24	9.74	7.66	8.26	8.13	8.63	8.63				
	THYRO1001800	6.25	3.20	2.10	5.72	6.37	4.12	5.27	7.47	7.47				
	THYRO1001803	19.67	13.46	12.81	14.42	14.10	16.43	18.72	18.13	18.13				
	THYRO1001809	3.63	3.46	2.67	3.26	4.76	4.46	2.91	5.54	5.54				
55	THYRO1001817	6.44	6.33	5.18	4.47	3.95	4.74	8.83	7.32	7.32*	-	*	+	
	THYRO1001819	5.55	5.75	5.06	8.24	5.95	6.83	5.79	6.86	6.86				

Table 340

5	THYRO1001120	3.6	3.56	2.97	4.01	3.89	3.81	3.24	4.59	4.59				
	THYRO1001121	4.68	3.13	2.03	5.64	4.07	3.90	2.70	4.05	4.05				
	THYRO1001128	6.11	5.32	3.34	12.06	12.42	10.51	5.36	6.39	6.39	**	+		
	THYRO1001133	6.15	4.73	4.57	9.2	11.55	7.92	6.41	7.28	7.28	*	+	*	+
	THYRO1001134	3.36	2.97	3.23	3.78	3.94	5.18	4.36	4.50	4.5		**	+	
	THYRO1001142	0.74	0.74	1.04	0.72	2.52	2.41	0.96	1.79	1.79				
10	THYRO1001173	15.19	9.02	12.22	26.91	29.74	31.51	28.83	31.54	31.54	**	+	**	+
	THYRO1001175	1.52	0.43	1.46	2.01	0.80	2.13	0.96	1.73	1.73				
	THYRO1001177	2.64	2.90	2.12	5.03	6.80	5.41	2.98	4.26	4.26	**	+		
	THYRO1001189	11.01	7.39	8.79	19.93	32.38	18.70	9.07	8.97	8.97	*	+		
	THYRO1001194	3.46	1.13	2.28	5.96	5.42	5.39	1.82	2.43	2.43	**	+		
15	THYRO1001204	4.45	2.95	2.30	6.96	6.86	8.50	3.26	4.79	4.79	**	+		
	THYRO1001205	24.03	16.88	15.68	32.39	32.90	31.15	22.06	24.66	24.66	**	+		
	THYRO1001213	3.76	2.34	2.06	5.73	8.42	6.51	4.19	4.49	4.49	*	+	*	+
	THYRO1001224	9.88	5.89	5.95	9.43	12.54	11.82	5.58	6.76	6.76				
	THYRO1001237	2.56	2.32	13.39	3.81	2.63	3.98	5.21	5.02	5.02		**	+	
20	THYRO1001242	27.87	23.01	22.93	21.64	25.67	32.15	25.14	28.77	28.77				
	THYRO1001258	3.57	5.51	4.92	4.9	6.74	6.73	7.47	5.30	5.3				
	THYRO1001262	1.72	1.10	1.83	6.36	5.01	5.41	2.24	3.79	3.79	**	+	*	+
	THYRO1001266	1.55	0.64	0.79	1.26	1.48	1.18	1.70	1.12	1.12				
	THYRO1001271	3.44	2.05	1.29	2.26	3.55	2.36	3.05	2.35	2.35				
	THYRO1001287	3.96	1.21	1.37	3.53	2.40	2.74	3.19	2.91	2.91				
25	THYRO1001290	1.14	0.69	1.23	1.44	2.26	2.04	2.54	3.09	3.09	*	+	**	+
	THYRO1001291	1.66	1.74	1.06	3.35	4.38	3.14	2.28	4.20	4.2	**	+	*	+
	THYRO1001297	5.89	5.62	3.44	7.28	6.73	6.27	3.04	3.57	3.57				
	THYRO1001302	0.7	1.17	1.36	2.14	3.01	3.14	1.40	2.26	2.26	*	+		
	THYRO1001313	4.31	2.12	1.72	3.28	3.86	2.48	2.67	3.67	3.67				
30	THYRO1001320	4.07	2.24	2.43	7.21	7.25	7.12	3.37	4.30	4.3	**	+		
	THYRO1001321	4.3	1.74	1.67	5.83	6.09	3.75	2.97	2.21	2.21				
	THYRO1001322	2.79	2.55	2.39	3.89	5.05	3.82	2.48	1.98	1.98	*	+		
	THYRO1001327	1.5	1.06	0.78	3.17	2.62	2.46	1.64	1.54	1.54	**	+		
	THYRO1001336	5.87	4.46	7.00	13.05	17.27	14.64	6.39	6.28	6.28	**	+		
35	THYRO1001347	0.03	0.55	0.25	0.69	2.15	0.73	1.35	0.54	0.54				
	THYRO1001358	11.06	9.93	9.25	14.71	16.38	14.53	9.85	8.62	8.62	**	+		
	THYRO1001363	5.86	3.17	4.11	5.35	3.91	6.10	4.52	5.65	5.65				
	THYRO1001365	5.19	2.07	3.95	4.26	3.12	4.83	2.55	3.93	3.93				
	THYRO1001374	9.65	2.81	3.50	6.43	5.39	7.37	3.94	7.65	7.65				
	THYRO1001401	7.01	3.08	4.71	9.44	10.37	11.91	6.83	6.19	6.19	*	+		
40	THYRO1001403	5.97	2.05	2.57	7.36	6.46	7.19	3.33	5.45	5.45				
	THYRO1001405	5.97	3.44	4.77	7.32	6.00	9.69	6.01	5.53	5.53				
	THYRO1001406	18.99	10.90	12.10	23.76	22.00	31.87	17.99	23.95	23.95	*	+		
	THYRO1001411	13.78	6.66	6.31	15.64	15.28	13.18	9.35	10.33	10.33				
	THYRO1001420	16.57	7.72	7.86	12.67	12.64	10.93	13.35	14.42	14.42				
45	THYRO1001426	12.94	7.75	6.41	21.71	18.55	24.48	11.12	13.81	13.81	**	+		
	THYRO1001430	8.77	5.32	6.79	6.79	9.85	9.38	6.22	8.03	8.03				
	THYRO1001434	4.36	1.78	2.26	3.34	4.87	4.04	2.27	4.13	4.13				
	THYRO1001456	6.47	2.68	3.34	4.42	3.96	4.89	4.38	4.91	4.91				
	THYRO1001457	6.96	3.92	4.84	6.7	6.42	9.36	5.12	7.08	7.08				
	THYRO1001458	9.57	4.98	6.73	8.45	4.79	10.94	7.33	10.09	10.09				
50	THYRO1001459	11.09	4.54	5.24	11.67	15.21	11.24	5.79	8.41	8.41				
	THYRO1001471	6.36	3.07	1.95	5.35	3.31	4.14	2.83	4.03	4.03				
	THYRO1001478	6.87	2.62	2.63	3.98	3.73	5.94	4.03	6.74	6.74				
	THYRO1001480	13.1	8.34	8.71	20.72	21.69	22.38	10.45	33.77	33.77	**	+		
	THYRO1001481	5.7	2.94	3.90	7.61	6.97	7.30	4.44	6.00	6	*	+		
55	THYRO1001487	7.46	5.22	5.93	9.3	8.06	9.99	6.17	7.84	7.84	*	+		
	THYRO1001495	11.89	6.81	10.31	8.41	6.19	9.91	4.61	6.76	6.76				

Table 339

5	THYRO1000783	1.63	0.89	1.30	3.1	2.26	1.53	1.80	1.30	1.3				
	THYRO1000786	4.89	2.61	2.30	6.28	3.05	5.87	4.15	4.10	4.1				
	THYRO1000787	10.6	5.80	4.42	7.07	6.40	5.00	7.52	6.30	6.3				
	THYRO1000792	6.58	1.87	1.67	2.34	3.23	1.91	2.22	2.34	2.34				
	THYRO1000793	2.04	0.81	0.90	2.24	3.46	2.95	1.63	1.90	1.9*	+			
	THYRO1000795	2.76	1.16	1.46	2.99	2.52	3.49	2.58	3.17	3.17				
10	THYRO1000796	2.38	0.64	1.44	4.8	3.84	4.16	2.52	2.59	2.59**	+			
	THYRO1000798	3.16	1.83	2.57	4.6	3.74	3.94	2.76	3.06	3.06*	+			
	THYRO1000800	7.44	4.89	4.90	15.05	11.25	16.69	6.56	6.96	6.96**	+			
	THYRO1000805	0.7	1.04	0.84	1.39	1.41	1.19	1.16	1.27	1.27*	+	*	+	
	THYRO1000815	7	4.02	3.01	10.69	12.71	10.92	7.46	5.49	5.49**	+			
15	THYRO1000829	4.85	1.50	0.99	3.49	4.27	2.08	2.62	2.36	2.36				
	THYRO1000835	2.11	1.21	1.15	2.86	3.23	3.63	2.50	4.32	4.32*	+	*	+	
	THYRO1000843	5.05	2.38	2.97	4.77	5.02	6.46	4.36	3.37	3.37				
	THYRO1000846	2.51	1.06	0.98	2.34	1.74	1.56	2.17	1.43	1.43				
	THYRO1000852	2.42	0.77	2.13	2.03	1.40	2.69	3.08	3.10	3.1				
20	THYRO1000855	4.5	4.43	3.85	5.88	4.56	7.12	5.76	3.18	3.18				
	THYRO1000865	3.16	2.10	3.34	4.86	6.09	6.43	5.14	2.65	2.65**	+			
	THYRO1000866	11.62	9.40	6.30	9.67	9.65	5.08	11.39	9.54	9.54				
	THYRO1000881	36.03	18.32	15.54	24.61	23.19	29.23	22.14	28.98	28.98				
	THYRO1000894	3.99	1.72	1.92	2.01	2.07	2.23	2.83	2.03	2.03				
25	THYRO1000895	2.03	0.86	1.43	1.55	2.22	2.83	1.11	1.40	1.4				
	THYRO1000916	3.35	1.86	1.68	6.43	4.60	5.32	3.15	2.84	2.84*	+			
	THYRO1000917	19.78	13.58	15.27	18.14	13.63	19.91	15.55	24.10	24.1				
	THYRO1000926	3.79	1.84	2.71	4.53	2.38	2.98	3.39	2.18	2.18				
	THYRO1000934	0.9	1.09	0.59	2.64	2.45	2.04	2.64	2.12	2.12**	+	**	+	
30	THYRO1000951	4.53	2.89	1.88	3.09	4.97	2.59	3.91	3.92	3.92				
	THYRO1000952	3.27	1.18	1.32	2.44	2.17	2.23	1.41	2.31	2.31				
	THYRO1000956	2.11	1.50	1.47	2.05	2.05	1.60	2.11	2.25	2.25				
	THYRO1000960	5.02	0.63	1.57	3.83	4.64	3.41	3.77	4.16	4.16				
	THYRO1000961	1.21	1.05	0.73	2.4	1.40	1.52	2.97	2.62	2.62		**	+	
	THYRO1000964	2.36	2.00	1.45	3.05	2.41	3.11	3.20	2.63	2.63				
35	THYRO1000971	6.39	3.74	2.87	7.64	6.60	7.93	4.97	5.58	5.58				
	THYRO1000974	8.5	6.07	6.15	9.83	9.20	11.43	9.21	8.90	8.9*	+			
	THYRO1000975	6.08	2.45	2.54	7.25	6.73	7.67	5.66	3.65	3.65*	+			
	THYRO1000983	6.75	2.78	2.84	5.03	3.45	3.63	5.16	7.50	7.5				
	THYRO1000984	4.73	2.02	2.56	6.84	6.78	4.19	3.85	4.94	4.94				
40	THYRO1000988	5.73	4.61	2.66	9.09	5.83	6.82	5.38	4.73	4.73				
	THYRO1000991	5.53	2.99	3.68	7.73	4.24	7.53	5.28	4.92	4.92				
	THYRO1000999	1.49	2.18	1.52	3.22	3.15	4.39	2.64	2.87	2.87*	+	*	+	
	THYRO1001003	3.32	1.87	1.67	2.91	2.45	1.95	2.38	1.98	1.98				
	THYRO1001015	6.07	3.22	4.17	6.03	3.39	4.75	4.51	4.29	4.29				
45	THYRO1001016	5.47	1.00	0.49	0.81	2.15	1.07	3.41	1.14	1.14				
	THYRO1001022	4.57	1.75	1.46	2.49	2.37	2.27	3.16	2.69	2.69				
	THYRO1001031	7	3.67	3.54	7.94	8.10	9.10	7.42	6.69	6.69*	+			
	THYRO1001033	2.8	0.57	1.23	2.39	2.37	1.06	1.41	2.32	2.32				
	THYRO1001062	3.82	2.25	2.08	5.76	5.14	5.15	3.45	3.99	3.99*	+			
	THYRO1001063	2.69	1.60	2.09	4.12	3.13	4.17	2.95	2.51	2.51*	+			
50	THYRO1001071	0.69	1.53	1.22	0.98	1.08	0.43	1.16	1.21	1.21				
	THYRO1001080	5.05	2.34	2.74	5.3	3.96	5.08	3.55	4.04	4.04				
	THYRO1001093	3.71	2.05	1.76	6.8	6.51	4.95	3.07	3.82	3.82*	+			
	THYRO1001100	2.79	1.59	1.28	2.23	2.44	2.67	1.71	3.71	3.71				
	THYRO1001102	4.56	2.46	2.61	2.98	3.38	2.67	4.62	4.11	4.11				
55	THYRO1001104	7.28	6.54	6.58	7.94	7.41	6.48	4.57	5.35	5.35	**	-		
	THYRO1001109	2.63	2.02	1.30	2.32	2.09	1.60	2.52	1.80	1.8				
	THYRO1001113	1.05	0.71	0.52	0.95	1.64	0.74	2.24	3.05	3.05	**	+		

Table 338

5	THYRO1000395	4.25	2.93	1.91	4.03	3.11	3.93	4.17	2.18	2.18				
	THYRO1000400	4.41	1.20	1.12	2.44	2.11	3.30	1.51	2.67	2.67				
	THYRO1000401	5.78	2.72	2.22	4.86	5.69	4.69	3.46	3.98	3.98				
	THYRO1000407	2.85	1.30	0.87	2.33	1.72	1.87	2.55	3.06	3.06				
	THYRO1000420	6.84	3.72	3.92	6.3	4.99	6.57	4.27	4.92	4.92				
	THYRO1000438	3.47	2.61	5.10	3.55	4.73	5.14	3.74	2.32	2.32				
10	THYRO1000452	3.79	2.27	3.32	4.32	3.39	3.80	3.50	2.68	2.68				
	THYRO1000455	0.86	0.19	0.08	0.98	0.97	1.02	0.43	0.69	0.69				
	THYRO1000471	3.13	0.99	1.71	4.82	2.11	3.45	2.03	2.21	2.21				
	THYRO1000481	3.05	2.09	1.78	2.49	2.59	3.24	2.75	3.65	3.65				
	THYRO1000484	7.3	2.87	2.29	10.67	15.51	6.38	4.46	3.81	3.81				
15	THYRO1000488	1.1	0.92	1.15	1.45	1.81	1.35	2.24	2.38	2.38	*	+	**	+
	THYRO1000501	2.42	1.63	1.50	2.59	2.38	2.19	2.65	3.01	3.01		*		+
	THYRO1000502	1.72	1.26	1.14	1.06	1.74	2.09	1.25	1.88	1.88				
	THYRO1000505	1.86	1.15	0.80	1	1.66	1.13	1.93	1.49	1.49				
	THYRO1000535	3.34	1.94	12.04	4.99	3.71	3.63	10.07	9.11	9.11			**	+
20	THYRO1000556	3.48	3.02	2.08	3.02	2.21	3.79	3.38	3.27	3.27				
	THYRO1000558	2.31	1.23	1.10	1.93	1.95	2.49	2.30	1.39	1.39				
	THYRO1000569	37.42	23.06	26.88	28.52	31.17	30.05	27.41	43.25	43.25				
	THYRO1000570	3.86	2.04	1.70	2.58	2.40	4.33	2.86	3.78	3.78				
	THYRO1000572	2.15	0.94	1.24	2.2	1.78	1.73	2.48	3.26	3.26		*		+
	THYRO1000573	2.15	0.40	1.11	1.23	2.42	1.75	1.79	2.04	2.04				
25	THYRO1000577	1.28	1.14	0.64	1.15	1.13	1.55	1.85	1.41	1.41				
	THYRO1000580	5.42	3.17	3.10	6.46	6.34	9.14	4.00	4.26	4.26	*	+		
	THYRO1000584	2.72	2.07	1.38	2.78	3.98	3.94	2.67	3.22	3.22				
	THYRO1000585	2.25	1.51	1.61	5.52	5.02	4.69	3.92	4.40	4.4	**	+	**	+
	THYRO1000596	0.84	0.25	0.33	0.85	1.98	1.44	1.19	1.17	1.17		*		+
30	THYRO1000602	5.45	3.58	2.07	8.38	7.15	5.61	4.80	5.98	5.98				
	THYRO1000605	3.06	1.73	1.76	2.38	1.83	1.39	2.18	2.05	2.05				
	THYRO1000615	1.88	0.80	0.63	1.19	1.72	1.17	1.04	2.25	2.25				
	THYRO1000625	3.03	2.54	1.58	4.59	3.95	5.93	3.48	4.60	4.6	*	+	*	+
	THYRO1000636	2.66	2.57	2.75	6.51	3.94	8.33	4.69	4.10	4.1	*	+	**	+
35	THYRO1000637	1.23	0.82	0.65	1.88	1.42	1.92	2.10	1.39	1.39	*	+		
	THYRO1000641	1.4	0.60	1.08	0.89	1.31	1.56	1.11	0.84	0.84				
	THYRO1000657	3.65	3.07	3.41	3.91	3.79	3.12	1.96	2.62	2.62		*		-
	THYRO1000658	7.81	3.42	3.03	11.25	11.55	11.93	5.08	5.90	5.9	*	+		
	THYRO1000662	2.88	1.16	0.83	2.17	1.76	1.90	1.97	1.81	1.81				
	THYRO1000666	2.42	0.88	1.16	3.25	2.79	4.33	1.98	2.43	2.43	*	+		
40	THYRO1000676	2.32	1.10	0.52	2.88	3.21	3.68	3.68	2.15	2.15	*	+		
	THYRO1000678	-0.09	0.33	0.95	0.54	0.74	1.28	1.19	2.92	2.92		*		+
	THYRO1000684	1.03	2.45	1.63	3.34	3.15	3.52	4.80	2.39	2.39	*	+		
	THYRO1000694	2.71	3.51	4.23	5.53	5.34	4.52	4.35	3.80	3.8	*	+		
	THYRO1000699	15.82	15.18	11.44	15.15	15.90	16.09	15.44	10.86	10.86				
45	THYRO1000712	3.39	2.96	2.14	8.58	5.42	7.84	3.11	4.20	4.2	*	+		
	THYRO1000715	4.02	2.34	2.31	2.86	4.26	2.85	3.39	2.68	2.68				
	THYRO1000716	2.32	0.65	1.04	2.97	4.44	2.89	2.03	1.56	1.56	*	+		
	THYRO1000717	2.15	0.84	1.30	4.23	5.94	4.84	1.47	3.93	3.93	**	+		
	THYRO1000723	0.84	0.47	0.25	0.76	1.49	1.41	0.88	0.44	0.44				
50	THYRO1000734	0.78	0.54	0.33	1.36	1.43	0.84	0.50	0.83	0.83	*	+		
	THYRO1000748	0.59	2.46	1.89	4.51	7.18	3.35	2.76	2.25	2.25				
	THYRO1000755	6.84	4.25	3.30	14.94	19.03	9.44	6.39	7.81	7.81	*	+		
	THYRO1000756	3.41	1.39	1.44	2.12	2.71	3.18	1.98	2.77	2.77				
	THYRO1000776	1.32	1.08	1.00	2.41	2.21	1.88	2.52	1.74	1.74	**	+	*	+
55	THYRO1000777	2.84	1.43	1.39	4.03	2.36	3.89	2.12	2.28	2.28				
	THYRO1000779	0.67	0.56	0.16	1.05	0.79	0.60	0.44	0.25	0.25				
	THYRO1000782	3.17	1.32	2.40	4.64	3.68	4.39	4.70	5.63	5.63	*	+	**	+

Table 337

5	THYRO1000036	1.47	2.88	1.52	4.59	3.60	4.55	3.37	2.09	2.09	*	+		
	THYRO1000040	3.94	3.60	4.34	8.08	4.14	6.42	4.83	5.15	5.15		*		+
	THYRO1000061	5.94	3.66	2.97	4.84	6.31	5.19	4.68	3.52	3.52				
	THYRO1000067	15.2	9.77	9.78	10.78	13.27	13.10	11.70	12.59	12.59				
	THYRO1000070	6	3.76	5.68	6.21	9.32	7.75	5.41	6.34	6.34				
	THYRO1000072	2.94	1.82	1.84	5.83	8.39	3.32	2.14	2.54	2.54				
10	THYRO1000084	4.5	1.85	2.58	3.76	4.67	3.19	3.46	2.16	2.16				
	THYRO1000085	10.88	13.54	13.23	14.79	17.02	16.91	12.99	15.14	15.14	*	+		
	THYRO1000086	0.12	1.27	1.00	1.39	0.92	1.37	0.61	1.10	1.1				
	THYRO1000087	0.56	0.67	0.91	1.37	1.09	0.98	1.47	0.51	0.51	*	+		
	THYRO1000092	6	2.56	1.98	8.27	6.56	7.42	3.48	3.45	3.45	*	+		
15	THYRO1000093	1.44	1.12	0.93	2.32	0.88	2.13	1.21	1.43	1.43				
	THYRO1000099	5.17	1.21	1.50	4.31	3.36	5.55	2.12	3.30	3.3				
	THYRO1000107	2.2	0.53	1.13	2.82	7.80	4.79	2.15	2.47	2.47				
	THYRO1000111	1.83	0.33	0.78	2.31	3.19	3.86	1.66	1.58	1.58	*	+		
	THYRO1000121	3.44	1.10	1.03	3.02	3.40	6.52	2.38	1.76	1.76				
20	THYRO1000124	2.37	0.51	0.78	3.06	2.51	2.25	0.89	1.60	1.6				
	THYRO1000129	1.3	0.82	0.51	1.26	1.52	1.53	0.49	1.02	1.02				
	THYRO1000130	3.62	2.11	2.49	5.24	10.43	5.75	7.64	2.92	2.92				
	THYRO1000132	8.41	1.76	1.74	4.45	6.81	7.03	2.87	3.11	3.11				
	THYRO1000134	3.55	1.81	2.95	6.64	4.07	4.40	3.58	4.01	4.01				
25	THYRO1000144	13.82	5.38	3.94	8.01	7.60	7.93	4.07	4.00	4				
	THYRO1000155	2.5	0.51	0.58	1.49	1.11	0.97	0.55	1.08	1.08				
	THYRO1000156	1.89	1.44	0.82	2.61	2.67	3.19	1.97	1.97	1.97	*	+		
	THYRO1000163	3.98	1.47	3.15	9.1	7.23	11.51	7.86	4.19	4.19	*	+		
	THYRO1000173	2.9	2.72	1.68	4.44	4.27	4.08	1.67	3.61	3.61	**	+		
30	THYRO1000186	9.1	5.19	4.20	10.2	15.51	9.61	7.74	7.44	7.44				
	THYRO1000187	5.63	2.01	3.20	6.21	7.01	6.32	5.05	3.18	3.18				
	THYRO1000190	2.89	1.46	2.17	5.4	4.76	5.31	4.40	2.66	2.66	**	+		
	THYRO1000196	0.92	0.80	1.33	2.19	1.72	1.35	0.94	1.18	1.18				
	THYRO1000197	3.18	2.33	2.51	5.88	3.71	6.16	4.77	4.51	4.51	*	+	**	+
	THYRO1000199	3.03	1.48	1.85	2.3	1.87	3.05	2.39	2.56	2.56				
35	THYRO1000206	14.52	5.55	4.65	11.65	9.64	12.12	6.54	6.11	6.11				
	THYRO1000221	5.01	1.90	2.05	5.6	6.77	7.34	2.67	3.86	3.86	*	+		
	THYRO1000222	7.73	2.24	1.94	3.18	4.68	4.24	4.78	2.83	2.83				
	THYRO1000223	1.72	0.91	0.91	5.64	4.49	4.50	3.42	4.40	4.4	**	+	**	+
	THYRO1000241	3.26	1.56	2.99	5.29	5.78	7.35	4.01	4.55	4.55	*	+	*	+
40	THYRO1000242	6.01	2.48	2.81	8.74	10.47	5.58	3.38	6.54	6.54				
	THYRO1000246	2.49	0.94	1.13	2.44	2.95	2.72	4.13	4.49	4.49			**	+
	THYRO1000253	3.03	2.39	2.12	4	3.56	6.64	2.35	3.27	3.27				
	THYRO1000270	0.85	0.93	0.64	2.95	1.36	0.98	0.55	0.45	0.45		*		-
	THYRO1000279	2.19	0.22	0.27	0.43	1.67	1.01	0.46	0.46	0.46				
45	THYRO1000285	6.19	3.69	1.88	4.45	4.41	6.07	8.92	4.83	4.83				
	THYRO1000288	7.58	2.67	2.64	4.38	5.78	3.82	4.63	6.75	6.75				
	THYRO1000296	3.95	2.07	1.83	3.07	4.49	3.23	3.68	3.54	3.54				
	THYRO1000320	4.13	0.95	0.96	3.75	3.95	6.99	3.33	5.20	5.2				
50	THYRO1000322	38.05	21.86	30.50	21.36	20.13	23.75	18.89	19.42	19.42				
	THYRO1000327	1.02	0.47	0.74	3.44	1.87	3.51	2.40	2.03	2.03	*	+	**	+
	THYRO1000343	3.18	0.96	1.50	2.27	1.34	2.27	1.96	1.19	1.19				
	THYRO1000345	4.6	2.12	2.05	3.98	5.40	3.79	1.33	1.91	1.91				
	THYRO1000358	7.71	5.28	3.61	7.26	4.42	4.45	5.44	7.71	7.71				
	THYRO1000368	11.25	3.81	3.69	7.91	6.70	6.11	5.37	4.82	4.82				
	THYRO1000375	6.52	5.33	3.32	11.74	11.72	9.07	7.23	13.34	13.34	*	+	*	+
55	THYRO1000381	1.08	0.73	0.85	2.03	1.76	1.73	1.91	1.07	1.07	**	+		
	THYRO1000387	2.85	2.46	2.45	4.71	4.58	5.18	3.81	2.92	2.92	**	+		
	THYRO1000394	3.11	2.36	2.61	4.86	4.51	5.33	6.21	6.15	6.15	**	+	**	+

Table 336

5	SKNMC1000099	4.27	1.82	4.32	2.68	2.85	4.25	4.90	2.31	2.31				
	SKNMC1000104	2.88	1.34	1.64	2.26	2.75	3.25	1.82	2.06	2.06				
	SKNMC1000113	2.91	1.98	1.70	2.53	3.12	2.50	2.17	2.08	2.08				
	SKNMC1000119	4.61	2.84	2.09	3.6	4.44	4.19	3.90	3.35	3.35				
	SKNMC1000142	2.86	0.96	0.73	2.73	1.96	2.31	2.39	2.51	2.51				
	SKNMC1000170	4.02	1.58	1.54	3.23	3.13	3.75	2.53	3.66	3.66				
10	SKNMC1000178	5.92	3.14	3.92	5.65	4.47	6.23	4.68	4.57	4.57				
	SKNMC1000194	3.57	2.37	1.14	2.02	1.84	1.46	1.82	1.68	1.68				
	SKNMC1000198	4.86	3.19	3.66	3.95	2.35	5.30	3.50	3.61	3.61				
	SKNMC1000225	3.86	1.48	1.25	3.04	2.83	3.41	1.69	1.50	1.5				
	SKNMC1000249	2.6	1.16	0.14	2.11	0.98	1.05	0.97	1.03	1.03				
15	SPLEN1000007	3.1	1.45	1.01	2.61	2.77	3.19	1.50	2.71	2.71				
	SPLEN1000012	4.58	1.70	1.35	3.53	2.59	2.41	3.41	4.25	4.25				
	SPLEN1000014	6.11	2.53	3.00	5.55	7.51	4.48	3.02	3.02	3.02				
	SPLEN1000036	2.67	1.59	1.60	2.81	3.21	2.90	3.30	2.69	2.69				
	SPLEN1000059	0.04	0.28	0.35	0.37	0.20	0.93	0.51	0.65	0.65		*		+
20	SPLEN1000068	2.47	1.01	1.48	3.14	3.20	4.62	4.16	2.46	2.46	*	+		
	SPLEN1000072	3.94	2.95	2.34	4.26	4.36	3.28	3.61	3.41	3.41				
	SPLEN1000101	41.57	16.82	24.85	23.9	21.81	9.24	15.06	12.84	12.84				
	SPLEN1000108	3.06	1.50	1.01	2.01	2.01	1.31	1.57	2.16	2.16				
	SPLEN1000113	4.35	2.46	2.67	4.83	2.55	2.28	3.11	3.66	3.66				
25	SPLEN1000114	2.42	2.37	1.43	3.43	2.78	2.56	2.74	3.97	3.97		*		+
	SPLEN1000132	4.91	2.27	3.07	3.65	2.33	4.08	4.07	4.65	4.65				
	SPLEN1000135	4.83	1.59	3.15	4.45	2.38	2.83	5.59	5.94	5.94		*		+
	SPLEN1000136	4.48	3.01	2.79	7.59	5.71	8.15	9.03	12.90	12.9	*	+	**	+
	SPLEN1000141	2.18	1.15	1.72	2.22	2.60	2.27	2.35	1.59	1.59				
30	SPLEN1000164	4.46	1.47	1.76	5.13	4.33	4.86	3.29	5.58	5.58				
	SPLEN1000166	2.49	0.67	1.05	2.36	3.89	2.42	2.08	3.68	3.68				
	SPLEN1000175	5.45	3.05	4.54	4.81	4.46	4.23	3.32	5.47	5.47				
	SPLEN1000182	2.6	0.65	0.61	1.52	1.41	2.22	1.31	1.69	1.69				
	SPLEN1000185	3.66	1.87	1.77	5.3	4.71	4.35	5.29	7.02	7.02	*	+	**	+
35	THYMU1000004	14.86	7.77	9.02	24.57	18.18	21.23	10.89	18.76	18.76	*	+		
	THYMU1000009	8.45	5.32	5.87	7.04	5.33	4.60	6.33	5.23	5.23				
	THYMU1000015	26.6	19.78	21.97	16.29	13.38	16.01	9.72	8.42	8.42	*	-	**	-
	THYMU1000016	8.26	4.04	3.89	15.26	18.83	11.55	9.39	7.02	7.02	*	+		
	THYMU1000023	3.89	1.34	1.23	2.77	2.08	3.06	2.39	2.39	2.39				
	THYMU1000034	2.61	1.47	0.66	2.74	1.63	1.39	1.31	3.64	3.64				
	THYMU1000035	1.07	0.61	0.61	0.44	0.64	0.76	1.85	2.01	2.01		**		+
40	THYMU1000037	1.82	1.82	1.19	2.22	2.35	0.98	2.22	2.11	2.11				
	THYMU1000042	10.49	6.31	8.55	6.35	4.98	6.18	8.88	5.36	5.36				
	THYMU1000047	4.11	2.46	3.11	10.3	9.57	11.11	4.37	4.74	4.74	**	+	*	+
	THYMU1000080	3.32	3.11	1.09	3.11	4.52	4.74	2.28	1.83	1.83				
45	THYMU1000094	32.63	25.01	18.12	54.59	42.21	15.66	23.80	19.03	19.03				
	THYMU1000109	8.44	4.34	3.79	6.74	8.15	5.93	7.23	6.42	6.42				
	THYMU1000127	6.78	3.40	3.18	8.92	8.62	7.88	6.21	6.83	6.83	*	+		
	THYMU1000130	4.13	1.20	1.02	4.32	4.36	3.32	2.41	3.21	3.21				
	THYMU1000137	4.62	2.65	2.71	3.35	5.77	3.60	4.29	4.56	4.56				
	THYMU1000146	4.71	3.58	4.49	7.3	4.71	6.30	5.63	4.56	4.56				
50	THYMU1000159	26.83	22.19	24.81	10.37	8.83	13.56	14.02	11.77	11.77	**	-	**	-
	THYMU1000163	6.99	6.16	7.74	9.39	10.76	10.43	5.92	8.13	8.13	**	+		
	THYMU1000167	2.34	1.29	1.70	2.93	3.51	2.43	1.61	1.52	1.52				
	THYMU1000186	5.07	2.17	2.10	3.12	3.14	2.53	2.77	2.94	2.94				
	THYRO1000017	5.52	2.12	2.08	4.91	6.19	6.15	4.38	3.34	3.34				
55	THYRO1000026	3.58	2.32	1.61	2.83	7.78	2.86	6.59	2.67	2.67				
	THYRO1000034	3.17	3.08	1.76	3.93	4.73	4.25	3.60	4.07	4.07	*	+		
	THYRO1000035	1.48	0.66	0.72	1.53	1.40	2.82	0.67	2.32	2.32				

Table 335

5	PLACE4000424	3.7	2.41	1.28	2.89	2.64	2.63	4.72	2.93	2.93				
	PLACE4000431	5.14	3.98	3.86	7.9	6.44	6.77	5.24	3.01	3.01	*	+		
	PLACE4000443	1.6	1.50	0.66	1.7	2.14	2.19	1.48	1.16	1.16				
	PLACE4000445	9.89	5.81	4.87	15.7	14.02	12.69	8.15	9.68	9.68	*	+		
	PLACE4000450	15.76	8.51	6.72	14.02	10.89	10.04	11.01	10.50	10.5				
10	PLACE4000455	3.87	3.67	2.19	8.55	5.76	6.75	4.27	7.65	7.65	*	+		
	PLACE4000465	6.69	5.73	3.42	9.19	8.96	7.57	6.23	7.71	7.71	*	+		
	PLACE4000466	31.49	24.03	27.55	30.7	30.16	27.24	58.59	49.41	49.41		**	+	
	PLACE4000472	17.06	12.16	12.26	19.04	18.92	24.52	24.99	19.96	19.96	*	+	*	+
	PLACE4000487	2.64	2.43	1.31	4.42	5.20	4.15	3.23	3.27	3.27	**	+		
	PLACE4000489	2.69	2.22	1.81	2.33	3.71	4.57	2.92	1.40	1.4				
15	PLACE4000494	6.6	3.79	3.88	6.95	7.91	8.87	5.80	5.92	5.92	*	+		
	PLACE4000502	21.16	12.73	11.94	19.98	23.69	17.79	12.36	16.13	16.13				
	PLACE4000521	6.7	5.05	4.78	4.05	6.11	3.01	4.55	6.40	6.4				
	PLACE4000522	4.91	3.07	3.08	7.26	9.24	7.69	9.03	9.77	9.77	**	+	**	+
	PLACE4000537	3.84	2.38	2.93	3.81	2.89	3.42	4.63	4.21	4.21		*	+	
20	PLACE4000548	2.58	1.71	3.60	3.4	2.67	4.50	1.35	2.28	2.28				
	PLACE4000558	0.39	0.54	0.56	2.25	2.45	2.36	1.46	1.14	1.14	**	+	**	+
	PLACE4000581	2.73	1.45	1.75	4.5	4.93	4.59	4.11	3.03	3.03	**	+		
	PLACE4000590	0.99	1.06	0.15	1.04	1.17	1.32	1.13	0.97	0.97				
	PLACE4000593	4.55	1.55	1.52	5.49	5.70	3.50	2.55	3.08	3.08				
25	PLACE4000612	14.51	9.28	7.13	10.09	12.95	7.67	9.14	12.79	12.79				
	PLACE4000638	3.93	2.21	3.37	3.98	5.06	3.32	3.69	4.06	4.06				
	PLACE4000650	1.03	1.91	1.53	2.69	2.70	2.58	3.71	1.90	1.9	*	+		
	PLACE4000651	8.37	7.37	5.41	16.13	16.91	20.29	11.75	11.67	11.67	**	+	**	+
	PLACE4000654	0.46	0.63	0.26	1.79	1.98	0.98	1.21	0.58	0.58	*	+		
30	PLACE4000670	1.04	0.70	1.13	2.04	2.89	1.47	0.74	0.43	0.43				
	PLACE4000685	23.26	12.26	10.49	28.55	27.61	40.89	20.42	24.20	24.2	*	+		
	PLACE4000687	0.45	0.07	0.48	0.48	0.65	1.00	0.21	0.78	0.78				
	PLACE5000003	2.7	1.36	1.81	2.51	2.87	2.69	2.63	1.48	1.48				
	PLACE5000005	2.1	1.91	0.92	1.98	1.29	2.16	2.69	3.30	3.3		*	+	
	PLACE5000019	1.64	0.35	0.54	1.85	0.86	1.29	2.04	1.56	1.56				
35	PLACE5000021	0.69	0.31	0.38	1.1	1.33	1.32	0.87	0.51	0.51	**	+		
	PLACE5000022	3.43	2.14	1.68	2.67	2.24	2.05	1.88	2.93	2.93				
	PLACE5000024	4.4	3.23	1.21	2.46	4.37	2.88	2.51	2.40	2.4				
	PLACE5000036	3.16	1.92	0.93	2.51	3.73	2.77	1.58	2.61	2.61				
	PLACE5000059	21.39	11.50	13.49	18.98	12.58	17.80	15.52	22.91	22.91				
40	PLACE5000076	1.04	0.14	0.59	0.44	1.09	3.27	1.13	0.58	0.58				
	PLACE5000117	6.61	3.04	3.55	6.57	7.00	6.53	6.39	6.85	6.85				
	PLACE5000143	6.9	3.66	5.74	7.55	3.91	6.50	6.13	5.78	5.78				
	PLACE5000152	1.01	0.83	0.51	1.68	1.63	1.58	1.45	0.95	0.95	**	+		
	PLACE5000154	2.82	2.00	1.84	2.88	1.96	2.91	1.39	2.76	2.76				
45	PLACE5000155	24.77	17.51	14.25	20.28	21.63	23.82	15.99	20.71	20.71				
	PLACE5000165	32.82	17.87	18.74	27.86	24.93	25.31	22.84	22.39	22.39				
	SKNMC1000004	6.53	6.43	3.51	11.48	11.51	13.01	5.92	10.64	10.64	**	+		
	SKNMC1000011	4.21	2.51	3.08	4.72	4.77	4.26	3.98	2.83	2.83				
	SKNMC1000013	2.24	1.08	1.20	1.57	0.87	2.15	1.79	2.08	2.08				
	SKNMC1000014	2.76	2.14	1.24	4.71	2.24	4.37	3.92	1.88	1.88				
50	SKNMC1000018	3.3	2.08	1.94	2.72	3.17	4.77	4.12	2.92	2.92				
	SKNMC1000020	4.56	2.73	1.89	3.66	1.77	4.25	2.81	2.80	2.8				
	SKNMC1000046	2.2	1.75	1.00	2.53	2.58	2.02	1.58	2.04	2.04				
	SKNMC1000050	2.33	0.87	1.04	1.57	3.28	2.53	3.94	4.06	4.06		**	+	
	SKNMC1000062	23.15	15.32	13.39	21.75	19.79	22.30	21.79	25.10	25.1				
55	SKNMC1000075	3.21	1.19	1.16	1.75	2.09	2.04	1.59	1.71	1.71				
	SKNMC1000082	5.24	2.03	1.78	3.5	2.07	2.24	2.55	1.68	1.68				
	SKNMC1000091	10.17	5.98	5.10	6.85	6.21	8.67	4.99	6.41	6.41				

Table 334

5	PLACE4000014	5.92	2.92	3.44	5.18	6.07	5.84	4.46	4.89	4.89				
	PLACE4000029	1.91	1.44	1.35	3.21	1.93	3.26	3.99	3.79	3.79			**	+
	PLACE4000034	2.6	1.30	1.44	3.92	3.82	4.60	4.01	3.41	3.41	**	+	*	+
	PLACE4000049	10.4	5.48	5.72	12.83	16.95	11.80	9.94	9.10	9.1	*	+		
	PLACE4000052	6.49	3.73	2.47	4.77	4.77	5.30	5.23	5.62	5.62				
	PLACE4000062	6.59	2.48	4.03	4.7	5.26	5.48	4.59	4.62	4.62				
10	PLACE4000063	7.7	3.50	3.52	6.91	6.71	9.08	5.77	5.40	5.4				
	PLACE4000089	2.96	1.45	2.33	5.97	4.11	5.63	4.54	4.57	4.57	*	+	**	+
	PLACE4000093	2.81	1.09	0.89	1.95	1.69	1.17	2.18	1.71	1.71				
	PLACE4000100	4.42	2.89	2.49	3.93	4.32	5.21	3.23	2.62	2.62				
	PLACE4000103	5.02	1.97	1.98	3.66	2.71	3.95	2.81	2.33	2.33				
15	PLACE4000106	8.72	4.11	3.74	4.38	5.75	4.55	4.28	4.16	4.16				
	PLACE4000128	7.39	4.68	3.31	9.85	9.72	8.43	7.44	6.38	6.38	*	+		
	PLACE4000129	6.04	2.07	2.84	4.76	6.70	6.24	4.40	2.79	2.79				
	PLACE4000131	8.08	5.12	4.57	12.93	9.62	6.75	8.38	9.08	9.08				
	PLACE4000147	1.54	0.95	0.56	0.28	1.32	1.44	1.32	1.12	1.12				
20	PLACE4000156	10.36	6.90	8.62	23.53	13.89	24.29	10.09	14.64	14.64	*	+		
	PLACE4000175	2.77	1.36	1.67	3	2.23	3.75	2.99	2.63	2.63				
	PLACE4000190	25.73	14.17	16.07	19.71	16.55	18.77	20.04	22.67	22.67				
	PLACE4000192	19.18	10.59	8.86	17.39	19.36	14.48	12.50	10.81	10.81				
	PLACE4000206	26.35	11.24	12.17	18.68	19.88	13.96	10.44	9.28	9.28				
25	PLACE4000211	17.59	9.35	9.22	14.45	14.14	14.09	11.01	11.86	11.86				
	PLACE4000214	3.16	2.15	2.41	4.6	3.22	2.93	3.58	2.23	2.23				
	PLACE4000222	5.13	3.77	3.41	7.67	6.23	6.64	5.04	5.14	5.14	*	+		
	PLACE4000223	5.15	2.40	3.83	4.77	3.40	3.75	4.17	5.28	5.28				
	PLACE4000229	2.61	1.29	1.59	3.13	1.82	2.66	3.16	3.28	3.28		*	+	
	PLACE4000230	10.54	4.47	5.13	3.92	4.50	6.23	2.12	1.74	1.74				
30	PLACE4000233	7.43	4.11	1.84	9.98	7.86	6.99	4.69	5.82	5.82				
	PLACE4000239	10.37	3.20	3.64	8.75	7.61	7.98	4.24	5.32	5.32				
	PLACE4000247	3.98	2.15	1.70	4.78	4.11	3.53	4.31	3.20	3.2				
	PLACE4000250	6.06	3.58	4.71	8.33	8.43	6.31	5.56	7.08	7.08	*	+		
	PLACE4000252	2.91	1.12	1.52	2.79	1.94	3.45	2.33	2.20	2.2				
35	PLACE4000259	8.04	3.19	7.29	6.61	5.24	7.03	5.35	5.02	5.02				
	PLACE4000261	12.86	7.43	11.27	7.94	6.30	11.29	13.49	12.71	12.71				
	PLACE4000264	5.07	2.86	1.88	6.35	6.52	5.02	3.87	4.16	4.16				
	PLACE4000269	8.57	4.36	5.52	8.01	9.34	7.35	6.12	5.77	5.77				
	PLACE4000270	3.13	1.82	0.87	2.42	1.82	3.08	1.61	2.16	2.16				
40	PLACE4000281	19.68	7.73	9.21	20.75	31.26	26.50	19.08	19.52	19.52	*	+		
	PLACE4000300	6.08	3.69	2.60	7.08	6.91	5.29	4.32	5.19	5.19				
	PLACE4000320	5.62	3.77	3.47	7.13	6.02	6.80	4.81	4.30	4.3	*	+		
	PLACE4000323	8.19	5.61	3.78	9.71	7.40	10.97	6.79	7.01	7.01				
	PLACE4000326	4.48	1.87	1.75	4.11	3.23	4.42	3.33	2.91	2.91				
	PLACE4000344	2.79	2.15	2.50	2.98	1.69	2.74	1.96	2.31	2.31				
45	PLACE4000347	20.7	10.82	8.58	19.27	12.61	11.57	8.40	11.08	11.08				
	PLACE4000354	4.74	1.02	1.75	4.04	3.76	1.42	1.52	3.10	3.1				
	PLACE4000367	2.52	1.53	1.38	2.65	2.91	2.67	2.13	2.39	2.39				
	PLACE4000369	4.83	2.57	3.31	5.06	4.32	4.54	4.37	5.39	5.39				
	PLACE4000379	5.69	3.46	3.45	6.11	6.24	7.35	4.30	5.61	5.61	*	+		
50	PLACE4000387	3.69	1.95	1.27	2.17	3.32	3.40	2.28	3.13	3.13				
	PLACE4000392	1.14	0.56	0.17	0.91	1.44	1.78	1.07	1.00	1				
	PLACE4000399	23.89	17.50	15.29	24.02	23.23	28.14	19.33	22.84	22.84				
	PLACE4000401	1.48	0.84	0.45	2.03	4.11	1.07	1.24	1.78	1.78				
	PLACE4000403	9.89	5.20	5.81	9.29	8.13	6.25	5.57	7.91	7.91				
55	PLACE4000411	5.72	2.12	2.75	5.81	4.30	5.15	4.86	3.29	3.29				
	PLACE4000415	3.21	2.22	2.69	3.67	4.43	3.44	4.57	6.28	6.28		**	+	
	PLACE4000416	4.63	3.13	2.08	5.57	4.05	4.56	5.50	4.53	4.53				

Table 333

5	PLACE3000199	3.29	1.08	1.38	2.04	1.59	1.81	1.36	3.52	3.52				
	PLACE3000205	9.93	4.59	5.70	17.83	17.57	18.45	14.66	13.74	13.74	**	+	*	+
	PLACE3000207	5.7	3.47	2.72	7.85	6.73	9.27	4.82	3.93	3.93	*	+		
	PLACE3000208	5.91	3.83	2.56	4.66	4.50	5.84	3.33	5.31	5.31				
	PLACE3000213	3.26	1.41	0.88	1.85	1.88	1.34	1.39	1.20	1.2				
	PLACE3000215	5.27	3.36	2.05	2.91	1.77	2.17	4.16	5.65	5.65				
10	PLACE3000218	0.67	1.20	0.52	0.53	0.72	1.11	0.94	1.60	1.6				
	PLACE3000220	4.81	2.27	2.38	5.89	5.17	5.82	4.14	4.16	4.16	*	+		
	PLACE3000221	18.58	12.33	11.49	19.49	17.73	21.75	11.62	11.46	11.46				
	PLACE3000225	2.26	1.52	1.43	2.24	4.06	3.45	1.47	2.45	2.45				
	PLACE3000226	4.27	2.49	2.02	2.27	5.71	4.75	1.91	2.73	2.73				
15	PLACE3000230	2.53	2.38	1.81	1.66	1.64	1.71	2.48	1.35	1.35				
	PLACE3000231	3.29	1.13	0.60	2.47	2.81	2.21	3.05	2.05	2.05				
	PLACE3000235	3.68	1.67	2.09	7.18	5.86	5.62	2.96	4.70	4.7	**	+		
	PLACE3000242	4.95	3.58	3.28	11.36	12.88	9.51	10.16	9.35	9.35	**	+	**	+
	PLACE3000244	1.78	1.29	0.91	1.71	1.41	0.91	1.35	0.85	0.85				
20	PLACE3000253	1.86	1.24	1.41	3.62	2.97	3.37	3.19	2.28	2.28	**	+	*	+
	PLACE3000254	51.54	34.63	40.51	40.03	46.12	56.93	50.43	47.16	47.16				
	PLACE3000271	5.35	3.90	4.49	12.52	15.43	16.28	5.75	8.41	8.41	**	+	*	+
	PLACE3000276	1.34	1.63	0.94	1.51	1.84	1.69	1.54	1.70	1.7				
	PLACE3000304	29.17	18.78	18.07	35.22	34.12	39.27	19.90	28.29	28.29	*	+		
25	PLACE3000309	5.85	2.02	1.54	4.32	5.65	5.33	3.03	4.10	4.1				
	PLACE3000310	2.86	0.49	0.75	1.95	1.51	1.29	0.96	1.26	1.26				
	PLACE3000320	2.43	0.72	1.39	2.35	2.67	2.63	2.57	2.39	2.39				
	PLACE3000322	3.17	2.14	2.01	4.49	4.42	5.13	3.26	3.42	3.42	**	+		
	PLACE3000330	3.98	4.24	5.26	4.75	5.64	8.32	9.28	8.32	8.32			**	+
	PLACE3000331	3.82	3.74	4.92	7.37	8.26	9.30	3.96	4.94	4.94	**	+		
30	PLACE3000336	2.26	2.25	2.90	3.09	4.08	3.48	1.74	3.42	3.42	*	+		
	PLACE3000339	1.51	1.25	0.97	2.83	3.03	1.44	3.34	1.37	1.37				
	PLACE3000341	4.76	1.28	2.07	6.03	6.07	5.79	3.01	2.61	2.61	*	+		
	PLACE3000350	3.67	2.80	1.30	3.28	4.94	3.47	3.39	2.10	2.1				
	PLACE3000352	6.03	5.05	2.30	5	5.48	4.50	3.98	4.70	4.7				
35	PLACE3000353	0.84	1.44	1.91	1.76	2.65	2.70	3.03	3.61	3.61			**	+
	PLACE3000362	1.98	1.66	1.84	6.16	5.62	6.95	2.53	2.39	2.39	**	+	**	+
	PLACE3000363	0.72	2.27	1.87	2.22	2.71	1.75	1.29	1.32	1.32				
	PLACE3000365	2.24	1.70	1.83	4.68	5.39	5.89	3.33	4.21	4.21	**	+	**	+
	PLACE3000373	1.03	0.43	0.22	0.96	1.41	0.94	0.42	0.29	0.29				
40	PLACE3000374	5.08	1.81	1.87	6.16	6.24	4.44	2.12	2.74	2.74				
	PLACE3000387	1.31	0.25	0.08	1.67	1.05	0.55	1.33	0.79	0.79				
	PLACE3000388	2.58	0.80	0.83	3.55	3.56	3.31	2.70	1.73	1.73	*	+		
	PLACE3000399	9.22	8.21	6.43	14.93	15.70	16.79	8.93	10.00	10	**	+		
	PLACE3000400	1.92	1.54	0.91	6.92	3.60	4.30	2.99	2.65	2.65	*	+	*	+
45	PLACE3000401	29	26.24	24.78	59.59	55.01	78.12	29.62	31.31	31.31	**	+	*	+
	PLACE3000402	2.02	1.57	1.10	4.22	3.77	2.97	1.86	1.95	1.95	**	+		
	PLACE3000405	6.4	2.32	4.16	6.78	5.01	5.58	4.43	5.58	5.58				
	PLACE3000406	4.28	1.49	2.84	5.5	4.66	5.13	2.47	2.85	2.85				
	PLACE3000413	8.22	3.55	3.63	4.09	5.81	4.91	5.48	4.88	4.88				
	PLACE3000416	4.22	2.84	2.70	5.29	3.87	4.91	3.53	2.90	2.9				
50	PLACE3000425	4.82	2.55	2.93	8.14	7.04	8.00	4.35	5.24	5.24	**	+		
	PLACE3000437	6.6	2.18	3.38	8.46	5.80	8.41	4.73	5.68	5.68				
	PLACE3000455	10.15	5.78	7.81	11.84	13.57	13.33	7.04	7.08	7.08	*	+		
	PLACE3000475	41.33	26.86	19.78	25.89	28.01	22.74	42.53	36.25	36.25				
	PLACE3000477	9.34	3.92	3.31	6.31	7.30	5.21	5.44	6.16	6.16				
55	PLACE4000003	2.47	1.31	0.94	1.5	2.41	1.58	1.49	1.63	1.63				
	PLACE4000008	5.72	2.63	3.64	8.57	12.47	10.37	7.46	7.86	7.86	*	+	*	+
	PLACE4000009	14.5	7.53	8.72	15.96	13.93	14.92	9.76	11.70	11.7				

Table 332

5	PLACE2000373	4.09	3.75	3.16	3.93	6.78	5.14	3.59	5.16	5.16				
	PLACE2000374	3.8	4.38	3.21	5.4	5.00	4.71	4.60	3.34	3.34	*	+		
	PLACE2000379	0.43	0.66	0.58	0.91	0.73	1.09	0.79	0.77	0.77	*	+	*	+
	PLACE2000386	263.51	193.15	186.41	112.96	134.53	97.90	242.44	237.17	237.2	*	-		
	PLACE2000388	6.14	2.57	3.20	4.18	4.37	4.11	3.57	5.67	5.67				
	PLACE2000392	22.7	12.68	10.22	19.04	26.24	23.82	20.84	18.58	18.58				
10	PLACE2000394	4.15	2.33	2.30	7.45	7.62	8.22	3.35	4.27	4.27	**	+		
	PLACE2000398	5.77	2.40	4.45	3.51	4.25	5.84	4.07	5.00	5				
	PLACE2000399	6.61	3.16	3.15	4.97	4.51	4.35	4.73	5.61	5.61				
	PLACE2000402	7.01	4.23	4.20	5.54	4.09	5.56	4.44	3.54	3.54				
	PLACE2000404	12.23	7.88	7.30	7.71	7.31	9.74	4.74	6.01	6.01				
15	PLACE2000411	21.27	11.68	11.82	11.14	10.88	25.73	14.78	18.35	18.35				
	PLACE2000418	5.51	3.37	3.01	6.69	5.87	6.09	4.87	3.75	3.75				
	PLACE2000419	7.28	4.27	3.30	7.57	9.49	8.40	4.83	4.59	4.59				
	PLACE2000425	4.32	2.24	3.29	5.08	4.37	6.06	3.45	3.86	3.86				
	PLACE2000427	6.26	3.55	3.23	4.54	4.54	5.08	5.10	5.28	5.28				
20	PLACE2000433	4.59	2.65	3.36	5.7	5.12	6.87	3.87	4.81	4.81	*	+		
	PLACE2000435	29.19	15.24	17.32	14.09	10.07	16.26	23.39	24.72	24.72				
	PLACE2000438	3.46	1.48	2.18	3.33	2.20	3.83	3.08	2.95	2.95				
	PLACE2000450	9.25	3.49	4.71	9.32	13.42	13.35	5.02	6.24	6.24	*	+		
	PLACE2000455	4.87	3.05	1.83	4.35	3.25	3.01	3.72	3.76	3.76				
25	PLACE2000458	7.14	3.76	3.85	4.27	6.42	5.62	5.42	5.04	5.04				
	PLACE2000464	10.07	4.31	6.99	6.94	8.11	6.92	5.43	8.55	8.55				
	PLACE2000465	5.73	2.78	3.87	8.13	9.58	9.56	5.26	6.47	6.47	**	+		
	PLACE2000473	17.94	8.98	12.76	32.72	23.26	29.31	35.66	50.78	50.78	*	+	**	+
	PLACE2000477	1.27	1.02	0.52	1.09	0.78	0.53	1.48	1.22	1.22				
30	PLACE3000004	7.55	3.19	4.53	8.79	7.24	9.45	5.46	5.75	5.75				
	PLACE3000009	61.9	29.47	28.32	32.27	25.30	29.38	45.27	58.28	58.28				
	PLACE3000020	9.44	5.05	5.57	6.59	7.39	6.52	4.82	4.55	4.55				
	PLACE3000029	9.17	4.67	4.83	9.55	12.07	7.65	6.59	5.44	5.44				
	PLACE3000038	3.05	1.65	1.71	3.75	5.45	4.67	2.86	3.09	3.09	*	+		
	PLACE3000052	4.37	2.71	2.77	5.23	4.15	6.64	3.13	2.24	2.24				
35	PLACE3000059	2.05	0.82	1.21	3.28	2.36	2.07	1.89	1.16	1.16				
	PLACE3000067	6.3	3.83	5.04	11.45	12.93	15.68	7.26	8.63	8.63	**	+	*	+
	PLACE3000069	5.9	3.04	3.53	5	5.11	8.56	5.67	5.68	5.68				
	PLACE3000070	27.81	15.78	20.14	32.22	26.02	53.33	21.90	29.50	29.5				
	PLACE3000103	2.43	0.95	1.30	3.54	4.94	4.26	1.89	2.90	2.9	*	+		
40	PLACE3000119	3.74	2.64	1.89	4.89	6.83	4.96	3.78	3.36	3.36	*	+		
	PLACE3000121	1.44	1.22	0.45	2.39	2.51	2.11	1.78	2.32	2.32	*	+	*	+
	PLACE3000124	5.32	4.70	4.50	12.73	13.74	11.54	6.57	8.87	8.87	**	+	*	+
	PLACE3000135	1.71	0.60	0.29	0.53	1.12	0.70	1.32	0.77	0.77				
	PLACE3000136	11.16	7.11	7.56	7.93	9.54	12.38	12.19	8.74	8.74				
	PLACE3000142	5.52	1.94	3.53	3.47	2.41	3.28	2.84	4.03	4.03				
45	PLACE3000145	6.76	3.35	3.37	7.36	7.49	6.80	6.17	8.06	8.06				
	PLACE3000147	10.95	4.59	5.26	7.55	8.27	7.90	4.11	4.34	4.34				
	PLACE3000148	2.39	0.34	0.58	0.98	1.42	1.44	1.27	2.88	2.88				
	PLACE3000154	1.15	0.41	0.41	0.66	1.09	1.15	0.77	2.42	2.42				
	PLACE3000155	7.02	4.85	4.71	7.95	8.38	9.70	5.71	8.16	8.16	*	+		
50	PLACE3000156	15.58	8.96	9.94	8.3	7.38	10.26	12.84	18.30	18.3				
	PLACE3000157	4.88	2.44	2.98	2.64	3.32	4.31	4.33	4.39	4.39				
	PLACE3000158	8.01	3.79	3.91	9.93	12.06	12.18	6.76	6.67	6.67	*	+		
	PLACE3000160	0.65	1.38	1.00	1.99	1.58	2.78	2.19	2.27	2.27		**	+	
	PLACE3000169	8.16	3.98	2.64	8.16	12.00	9.90	6.56	5.46	5.46				
55	PLACE3000181	10.19	6.03	5.74	6.99	7.06	6.27	8.77	10.50	10.5				
	PLACE3000194	3.17	1.81	1.47	3.38	2.68	2.61	2.86	3.96	3.96				
	PLACE3000197	0.71	0.07	1.20	1.17	0.86	1.14	4.41	2.01	2.01				

Table 331

5	PLACE2000039	6.48	4.35	4.61	11.87	11.66	13.79	6.80	7.28	7.28	**	+	*	+
	PLACE2000043	2.47	1.44	2.20	2.31	3.69	3.32	3.41	4.52	4.52			*	+
	PLACE2000044	5.02	3.35	3.46	5.51	3.83	5.89	4.93	7.31	7.31				
	PLACE2000047	8.18	4.36	3.83	9.19	11.31	14.75	5.33	7.74	7.74	*	+		
	PLACE2000050	12.24	3.78	3.08	8.61	10.29	7.90	7.32	6.64	6.64				
10	PLACE2000061	2.92	0.96	0.97	1.52	0.96	1.26	1.35	1.85	1.85				
	PLACE2000062	4.77	2.50	2.13	5.58	5.65	5.45	2.96	5.42	5.42	*	+		
	PLACE2000072	2.7	1.26	2.16	2.17	3.44	2.93	1.74	2.43	2.43				
	PLACE2000073	1.69	0.72	0.84	1.41	0.59	1.30	1.70	1.52	1.52				
	PLACE2000097	13.16	8.11	9.49	11.41	12.05	13.08	7.86	8.83	8.83				
15	PLACE2000100	5.14	3.46	2.83	5.96	4.13	5.86	4.27	5.06	5.06				
	PLACE2000103	4.64	3.10	3.20	7.22	5.44	6.13	4.03	3.95	3.95	*	+		
	PLACE2000106	7.76	2.85	4.06	6.8	7.28	7.13	4.31	4.99	4.99				
	PLACE2000111	4.84	2.29	3.47	5	5.26	5.57	4.32	7.27	7.27				
	PLACE2000115	2.29	0.90	1.18	1.38	0.91	1.85	2.19	2.02	2.02				
	PLACE2000118	40.98	28.15	29.38	32.74	33.08	38.40	30.44	42.97	42.97				
20	PLACE2000124	16.57	10.11	11.57	19.83	25.65	30.81	16.15	17.74	17.74	*	+		
	PLACE2000132	7.64	4.32	5.67	5.55	4.79	4.71	7.51	6.44	6.44				
	PLACE2000136	1.78	0.82	1.05	1.68	1.61	1.41	1.31	1.62	1.62				
	PLACE2000137	6.66	4.19	3.94	4.2	3.59	5.28	3.96	5.37	5.37				
	PLACE2000140	9.31	3.10	5.25	7.95	10.19	7.07	4.50	6.74	6.74				
25	PLACE2000147	2.32	1.00	0.75	2.39	2.55	2.14	1.33	2.93	2.93				
	PLACE2000153	1.79	0.33	0.76	0.89	1.36	1.15	2.17	2.54	2.54		*	+	
	PLACE2000164	2.92	1.24	1.74	1.97	2.41	1.94	1.21	2.25	2.25				
	PLACE2000170	4.49	2.57	2.11	5.8	5.33	5.19	3.14	3.80	3.8	*	+		
	PLACE2000172	3.21	1.40	2.70	1.1	3.14	2.28	1.52	1.72	1.72				
30	PLACE2000173	4.05	3.41	2.95	5.72	7.77	7.43	3.82	4.53	4.53	**	+		
	PLACE2000174	2.94	1.68	2.28	3.36	3.27	4.06	2.97	2.61	2.61	*	+		
	PLACE2000176	6.55	2.90	2.44	6.47	6.24	4.58	3.30	4.24	4.24				
	PLACE2000187	4.34	2.14	1.78	5.63	3.41	5.66	3.80	4.31	4.31				
	PLACE2000216	4.17	2.38	2.18	6.97	6.14	5.24	7.33	12.03	12.03	*	+	*	+
	PLACE2000219	5.75	2.86	2.79	6.33	5.19	5.66	5.15	5.03	5.03				
35	PLACE2000221	6	4.55	4.10	13.08	11.16	10.61	6.14	6.36	6.36	**	+		
	PLACE2000223	0.66	0.04	0.44	2.56	1.05	0.74	1.35	0.62	0.62				
	PLACE2000231	2.73	2.97	1.35	3.88	3.66	2.81	3.23	2.76	2.76				
	PLACE2000235	5.15	3.31	3.10	9.33	15.20	9.28	4.35	5.69	5.69	*	+		
	PLACE2000246	9.05	5.03	3.92	8.93	10.34	8.27	5.30	6.19	6.19				
40	PLACE2000244	4.4	2.75	1.21	7.23	5.72	5.03	3.18	4.43	4.43	*	+		
	PLACE2000274	8.27	4.14	5.09	4.88	4.54	3.46	4.83	6.06	6.06				
	PLACE2000287	14	9.69	10.03	11.98	14.31	14.19	12.42	12.37	12.37				
	PLACE2000296	3.51	1.96	2.07	2.61	2.73	3.24	2.29	3.69	3.69				
	PLACE2000302	2.31	2.23	2.10	3.57	4.89	5.77	3.81	3.32	3.32	*	+	**	+
	PLACE2000305	7.13	5.46	4.88	12.44	18.75	14.01	6.85	6.47	6.47	*	+		
45	PLACE2000317	1.79	1.81	1.59	2.18	2.88	3.79	2.56	2.49	2.49		**	+	
	PLACE2000324	1.64	0.45	0.66	1.23	0.90	1.15	1.56	1.03	1.03				
	PLACE2000334	4.7	3.19	3.38	3.36	3.85	3.08	3.51	4.53	4.53				
	PLACE2000335	6.89	3.67	3.94	9.95	12.98	11.87	4.72	8.12	8.12	**	+		
	PLACE2000340	1.92	1.00	1.25	2.13	2.37	2.25	1.13	1.70	1.7	*	+		
50	PLACE2000341	4.05	3.76	4.37	3.33	6.79	4.35	3.93	3.97	3.97				
	PLACE2000342	5.08	6.69	5.71	7.14	7.41	7.08	6.86	8.97	8.97	*	+	*	+
	PLACE2000347	4.37	5.20	4.34	7.13	9.07	10.11	6.32	7.54	7.54	*	+	**	+
	PLACE2000357	9.87	8.86	7.75	8.78	12.51	9.80	8.70	9.73	9.73				
	PLACE2000358	4.58	2.20	2.55	4.28	5.06	2.32	5.59	6.09	6.09		*	+	
55	PLACE2000359	2.5	0.52	0.79	3.39	1.81	2.56	2.27	0.82	0.82				
	PLACE2000366	6.64	3.37	2.29	8.44	9.84	6.44	5.08	3.45	3.45				
	PLACE2000371	4.65	3.72	1.76	1.73	2.65	2.33	2.69	2.16	2.16				

Table 330

5	PLACE1011675	0.49	0.41	0.33	1.54	2.62	1.74	1.03	0.63	0.63	**	+		
	PLACE1011682	1.44	1.27	1.77	1.27	1.81	1.50	0.94	2.14	2.14				
	PLACE1011708	4.35	4.02	4.14	5.7	7.61	8.08	4.28	4.88	4.88	*	+		
	PLACE1011719	1.76	1.55	1.39	2.03	3.35	3.13	2.09	2.66	2.66	*	+	*	+
	PLACE1011725	4.47	2.20	1.51	6.52	4.79	5.08	4.70	3.97	3.97				
10	PLACE1011729	2.26	0.34	1.16	2.9	3.70	2.58	1.88	1.07	1.07	*	+		
	PLACE1011741	1.85	1.08	1.46	2.17	2.55	1.44	1.47	2.04	2.04				
	PLACE1011749	4.07	1.97	2.35	5.14	5.66	5.55	2.94	2.96	2.96	*	+		
	PLACE1011757	7.95	5.78	4.73	28.51	35.97	33.70	18.45	19.91	19.91	**	+	**	+
	PLACE1011762	0.6	0.62	0.64	1.51	2.40	1.43	1.31	1.98	1.98	*	+	**	+
15	PLACE1011778	0.68	0.80	0.85	0.72	1.28	1.91	1.16	0.80	0.8				
	PLACE1011783	3.26	3.37	4.33	9.36	8.72	9.68	4.31	3.99	3.99	**	+		
	PLACE1011795	2.41	0.78	0.71	3.25	3.16	2.10	1.75	0.51	0.51				
	PLACE1011810	1.09	0.35	0.96	0.57	0.96	0.71	1.04	1.41	1.41				
	PLACE1011824	1.1	0.61	0.73	1.63	1.19	1.20	1.70	1.61	1.61		**	+	
	PLACE1011825	19.56	10.93	11.42	10.37	11.28	11.36	8.08	10.44	10.44				
20	PLACE1011835	2.12	1.20	1.49	1.76	1.50	1.06	1.49	0.95	0.95				
	PLACE1011836	32.53	15.61	18.36	27.63	35.75	28.68	27.23	20.95	20.95				
	PLACE1011847	0.74	1.05	0.62	0.62	0.87	1.11	1.60	1.10	1.1				
	PLACE1011853	1.16	0.16	0.77	0.69	0.71	1.38	0.70	1.13	1.13				
	PLACE1011858	2.38	2.07	1.60	2.19	2.08	1.60	2.79	2.84	2.84		*	+	
25	PLACE1011874	3.25	1.54	2.03	4.69	4.12	4.23	2.47	3.11	3.11	*	+		
	PLACE1011875	1.26	0.66	0.64	1.26	1.14	1.27	0.79	0.74	0.74				
	PLACE1011877	6.46	2.58	3.09	3.53	2.30	3.26	2.14	3.12	3.12				
	PLACE1011891	1.77	0.88	0.81	1.69	1.67	1.68	1.49	2.31	2.31				
	PLACE1011896	0.86	0.25	0.26	0.37	0.26	0.57	0.67	0.48	0.48				
	PLACE1011920	2.91	0.83	1.76	1.44	1.22	2.34	1.43	1.54	1.54				
30	PLACE1011922	4.71	2.40	2.11	4.92	2.79	4.42	3.68	4.23	4.23				
	PLACE1011923	3.63	1.24	1.28	5.32	2.65	2.76	7.49	10.90	10.9		**	+	
	PLACE1011937	6	2.51	3.82	3.74	4.24	5.24	4.33	4.96	4.96				
	PLACE1011939	4.24	2.12	2.87	2.83	3.92	4.33	4.29	5.83	5.83		*	+	
	PLACE1011940	5.02	1.82	3.30	7.08	7.36	8.48	4.28	5.85	5.85	*	+		
35	PLACE1011962	13.26	6.64	7.98	11.22	11.07	13.01	8.70	9.69	9.69				
	PLACE1011964	2.09	0.16	0.88	0.97	0.96	0.80	0.82	1.29	1.29				
	PLACE1011978	6.83	5.17	5.96	14.23	9.13	15.11	5.12	9.01	9.01	*	+		
	PLACE1011980	5.54	2.72	4.54	9.74	12.59	11.01	4.66	6.64	6.64	**	+		
	PLACE1011981	6.65	3.37	3.46	5.38	6.10	5.01	3.81	4.69	4.69				
40	PLACE1011982	0.91	0.32	0.06	0.49	1.81	1.02	0.79	1.02	1.02				
	PLACE1011995	4.44	2.50	2.12	5.89	6.15	5.90	3.97	3.81	3.81	*	+		
	PLACE1012023	1.79	0.70	1.25	1.43	1.61	2.10	1.24	1.17	1.17				
	PLACE1012026	1.87	0.19	0.62	1.01	0.13	0.38	0.66	0.81	0.81				
	PLACE1012031	2.22	1.02	2.34	1.31	1.21	3.28	1.23	2.49	2.49				
45	PLACE2000003	10.16	5.53	7.19	14.74	10.74	18.79	8.25	10.16	10.16				
	PLACE2000005	4.58	2.43	2.29	4.4	3.89	4.52	3.11	4.04	4.04				
	PLACE2000006	6.31	3.28	0.91	2.52	4.28	3.42	2.99	2.00	2				
	PLACE2000007	3.33	1.18	1.86	1.87	3.29	3.13	2.74	4.18	4.18				
	PLACE2000011	6.03	2.56	4.49	6.77	5.68	6.77	3.61	4.33	4.33				
	PLACE2000014	0.21	0.98	0.82	1.07	2.07	1.53	2.21	2.03	2.03		**	+	
50	PLACE2000015	1.83	0.65	1.01	1.76	1.29	2.19	1.52	0.85	0.85				
	PLACE2000017	3.21	0.73	1.74	6.58	4.80	4.12	2.77	1.87	1.87	*	+		
	PLACE2000021	3.22	1.34	2.09	3.94	3.51	4.88	1.82	2.10	2.1	*	+		
	PLACE2000022	7.75	3.64	2.82	8.01	9.76	8.90	3.63	6.09	6.09				
	PLACE2000030	8.7	4.13	5.99	6.21	7.35	6.10	5.75	6.25	6.25				
55	PLACE2000032	4.4	0.93	2.84	5.81	6.65	4.83	2.50	4.03	4.03				
	PLACE2000033	1.83	1.13	0.57	2.93	3.15	2.60	1.56	1.76	1.76	*	+		
	PLACE2000034	2.2	2.03	1.49	1.47	1.71	2.75	1.92	3.95	3.95				

Table 329

5	PLACE1011263	4.63	1.43	2.06	5.15	3.30	5.05	3.84	4.54	4.54				
	PLACE1011273	0.96	0.21	0.03	0.28	0.62	1.29	0.76	0.83	0.83				
	PLACE1011278	6.81	4.02	5.42	10.67	8.60	12.25	6.32	6.99	6.99	*	+		
	PLACE1011289	5.66	2.33	3.18	4.65	3.12	5.27	3.01	3.39	3.39				
	PLACE1011291	16.28	11.06	10.52	7.72	9.80	6.81	14.94	17.29	17.29				
10	PLACE1011296	3.24	2.37	2.66	4.3	4.86	3.54	3.68	3.04	3.04	*	+		
	PLACE1011310	4	1.37	1.23	4.91	7.48	2.45	2.90	2.71	2.71				
	PLACE1011311	6.86	4.63	5.58	11.54	13.47	10.02	8.99	6.21	6.21	**	+		
	PLACE1011321	2.48	2.00	2.29	4.17	3.53	4.74	3.10	3.06	3.06	**	+	**	+
	PLACE1011325	2.45	1.16	0.85	2.15	1.85	2.50	1.87	1.38	1.38				
15	PLACE1011332	2.06	1.37	1.10	2.9	1.77	3.23	1.54	3.88	3.88				
	PLACE1011340	4.71	2.86	3.96	6.93	7.43	10.39	3.26	4.42	4.42	*	+		
	PLACE1011353	8.94	8.02	6.47	12.12	12.68	8.45	5.57	6.13	6.13				
	PLACE1011360	5.26	2.74	2.31	7.14	13.29	6.66	11.83	17.54	17.54		**	+	
	PLACE1011364	3.45	2.09	2.62	4.62	3.01	2.44	3.75	3.95	3.95		*	+	
	PLACE1011365	2.35	1.17	0.95	2.03	1.96	2.41	0.96	2.66	2.66				
20	PLACE1011371	5.16	2.45	2.43	5.08	2.86	3.23	3.60	3.42	3.42				
	PLACE1011375	2.23	1.21	1.56	1.86	1.08	1.78	1.86	1.55	1.55				
	PLACE1011386	8.63	5.02	6.24	7.07	6.54	8.61	7.88	10.06	10.06				
	PLACE1011399	1.83	1.09	0.89	5.72	1.66	3.52	2.58	2.39	2.39		*	+	
	PLACE1011406	5.14	2.34	2.53	3.24	3.16	4.75	3.03	4.67	4.67				
25	PLACE1011407	5.6	2.12	1.49	5.65	6.78	4.60	3.19	4.91	4.91				
	PLACE1011419	3.79	1.50	2.18	3.71	3.80	3.26	2.85	4.10	4.1				
	PLACE1011433	3.79	3.19	4.12	13.24	18.92	14.07	5.04	9.50	9.5	**	+	*	+
	PLACE1011440	3.69	0.88	2.02	3.25	2.87	3.33	3.41	3.73	3.73				
	PLACE1011452	3.56	2.32	3.25	5.65	6.92	7.14	3.10	4.52	4.52	**	+		
30	PLACE1011465	1.9	0.93	1.60	1.74	1.90	2.00	2.17	2.04	2.04				
	PLACE1011472	5.01	1.93	2.18	2.83	4.34	2.95	3.24	2.62	2.62				
	PLACE1011477	7.19	3.67	4.99	9.17	8.71	7.03	6.34	7.80	7.8				
	PLACE1011478	4.7	2.46	2.21	8.34	7.12	6.47	4.52	4.33	4.33	*	+		
	PLACE1011492	5.64	3.42	3.03	6.13	7.41	5.44	6.73	7.31	7.31		*	+	
	PLACE1011498	2.62	0.69	0.77	2.57	1.98	3.73	3.99	6.38	6.38		*	+	
35	PLACE1011501	1.42	0.15	0.63	0.39	1.02	1.21	0.37	2.92	2.92				
	PLACE1011503	1.26	0.38	0.38	0.56	0.61	1.28	0.65	1.73	1.73				
	PLACE1011509	2.69	1.51	1.96	4.97	3.77	5.32	2.73	3.70	3.7	*	+	*	+
	PLACE1011514	3.56	3.02	2.30	6.22	6.65	9.32	5.42	3.34	3.34	*	+		
	PLACE1011516	9.2	5.83	6.52	8.39	11.75	7.44	6.58	7.31	7.31				
40	PLACE1011520	0.67	(0.02)	0.19	0.61	0.97	0.53	0.59	1.82	1.82				
	PLACE1011538	2.38	1.78	1.26	2.2	1.67	2.66	4.47	5.04	5.04		**	+	
	PLACE1011555	2.73	2.02	1.56	3.06	1.29	2.65	3.42	3.23	3.23		*	+	
	PLACE1011561	0.88	0.17	0.48	1.89	2.38	1.56	4.39	6.30	6.3	*	+	**	+
	PLACE1011563	3.61	1.68	1.69	2.85	2.58	2.68	3.94	2.74	2.74				
45	PLACE1011567	2.71	2.13	1.59	4.37	4.64	3.78	1.76	2.11	2.11	**	+		
	PLACE1011569	0.28	1.00	0.40	1.55	1.16	1.18	0.73	0.92	0.92	*	+		
	PLACE1011576	30.78	17.05	20.91	58.8	58.74	38.79	22.26	24.68	24.68	*	+		
	PLACE1011586	5.24	2.45	1.64	5.28	3.51	3.90	2.49	2.57	2.57				
	PLACE1011635	1.82	1.02	0.96	2.22	1.28	1.79	1.86	3.48	3.48				
	PLACE1011641	0.55	0.39	0.51	0.79	0.28	0.18	0.89	0.98	0.98		**	+	
50	PLACE1011642	2.33	1.95	2.07	3.44	2.17	2.75	1.72	3.63	3.63				
	PLACE1011643	1.74	0.86	1.81	2.9	2.61	2.81	1.78	2.26	2.26	*	+		
	PLACE1011646	4.54	1.91	2.30	4.88	5.46	7.17	3.47	4.39	4.39				
	PLACE1011649	5.04	2.68	5.39	5.34	8.26	6.45	8.02	7.02	7.02		*	+	
	PLACE1011650	9.82	9.33	4.23	8.72	9.82	8.25	7.64	7.21	7.21				
55	PLACE1011661	4.13	2.90	2.81	7.54	8.51	8.47	3.11	3.99	3.99	**	+		
	PLACE1011664	2.28	2.16	2.82	2.3	4.01	2.45	1.92	2.20	2.2				
	PLACE1011672	1.34	0.43	0.59	1.98	2.38	1.65	1.43	1.75	1.75	*	+	*	+

Table 328

5	PLACE1010877	3.9	0.68	2.81	4.57	8.26	6.30	4.12	5.68	5.68				
	PLACE1010882	1.73	0.87	-1.34	0.94	1.22	1.41	1.64	2.79	2.79				
	PLACE1010891	1.31	1.05	1.38	1.34	2.82	2.67	1.60	1.74	1.74		*	+	
	PLACE1010896	2.03	1.93	1.21	5.65	5.89	6.07	2.71	4.67	4.67	**	+	*	+
	PLACE1010900	7.45	5.19	4.52	6.71	10.28	6.75	5.29	6.78	6.78				
10	PLACE1010916	1.58	1.17	1.07	2.47	2.58	1.67	1.27	2.26	2.26	*	+		
	PLACE1010917	1.05	0.96	0.11	1.61	1.38	1.11	1.25	1.13	1.13				
	PLACE1010924	2.09	0.79	0.68	3.58	1.12	1.06	1.53	2.87	2.87				
	PLACE1010925	6.95	5.48	6.26	14.3	13.92	11.11	10.38	11.87	11.87	**	+	**	+
	PLACE1010926	4.68	2.80	3.56	5.61	3.87	4.95	5.17	4.94	4.94				
15	PLACE1010942	9.58	6.01	6.54	10.63	11.10	11.71	7.84	8.22	8.22	*	+		
	PLACE1010943	34.04	17.63	26.11	27.44	25.58	32.27	17.16	17.20	17.2				
	PLACE1010944	4.16	2.44	1.53	4.69	4.52	3.10	3.60	3.71	3.71				
	PLACE1010947	3	1.38	1.06	4.09	3.59	3.17	3.08	1.80	1.8				
	PLACE1010954	5.64	1.64	2.41	6.89	7.16	7.06	4.57	2.95	2.95	*	+		
	PLACE1010960	2.56	1.87	3.84	3.46	4.48	4.07	2.90	5.57	5.57				
20	PLACE1010965	2.32	1.81	1.90	3.82	3.17	4.63	3.08	3.88	3.88	*	+	**	+
	PLACE1010968	2.01	2.04	1.40	2.48	1.55	2.68	3.26	2.68	2.68		*	+	
	PLACE1010978	2.64	1.65	3.12	2.67	4.61	3.98	4.33	3.15	3.15				
	PLACE1010982	0.32	0.44	1.17	1.43	1.48	1.69	0.82	1.16	1.16	*	+		
	PLACE1010990	1.25	1.65	1.41	1.21	2.03	3.15	1.56	2.02	2.02				
25	PLACE1011017	4.02	2.33	2.07	4.93	5.02	3.31	2.67	2.53	2.53				
	PLACE1011019	4.19	3.51	2.69	3.28	4.24	3.10	3.40	4.43	4.43				
	PLACE1011026	0.53	0.56	0.94	1	1.14	1.01	1.44	1.90	1.9		**	+	
	PLACE1011032	1.04	1.26	1.14	1.35	3.76	1.41	1.45	1.09	1.09				
	PLACE1011041	2.22	2.15	1.83	3.19	3.33	2.65	2.20	2.59	2.59	*	+		
30	PLACE1011045	4.26	2.55	2.05	3.25	2.44	2.71	2.62	4.66	4.66				
	PLACE1011046	2.58	2.74	2.45	7.65	5.20	7.76	2.85	2.98	2.98	**	+	*	+
	PLACE1011054	5.53	5.97	3.21	7.9	9.44	10.30	5.46	7.19	7.19	*	+		
	PLACE1011056	12.06	8.95	6.62	12.16	14.46	16.30	10.02	8.27	8.27				
	PLACE1011057	1.87	1.37	1.37	4.52	4.54	3.70	1.84	1.48	1.48	**	+		
	PLACE1011059	0.6	0.44	0.37	1.23	1.32	1.03	0.88	0.52	0.52	**	+		
35	PLACE1011066	4.38	2.43	3.65	12.2	18.07	8.10	6.79	9.58	9.58	*	+	**	+
	PLACE1011067	8.25	6.00	13.05	11.16	14.93	13.07	8.67	11.65	11.65				
	PLACE1011090	3.34	3.04	3.20	4.44	8.79	7.02	2.18	1.69	1.69	*	+	**	-
	PLACE1011109	4.01	3.02	3.89	8.31	10.21	9.01	3.52	3.74	3.74	**	+		
	PLACE1011114	3.2	3.86	3.47	4.71	3.90	4.13	3.10	3.01	3.01				
40	PLACE1011116	10.05	5.20	4.98	6.55	11.37	7.07	9.74	10.38	10.38				
	PLACE1011122	1.51	0.61	0.83	0.61	2.18	1.37	1.66	1.51	1.51				
	PLACE1011133	3.84	1.08	1.97	3.52	5.23	3.54	3.50	2.83	2.83				
	PLACE1011134	3.94	1.65	2.34	3.61	3.61	3.94	2.95	3.46	3.46				
	PLACE1011143	3.34	0.78	1.07	1.6	0.98	1.60	1.65	1.94	1.94				
45	PLACE1011146	5.79	3.61	4.24	4.94	3.96	4.87	4.89	7.01	7.01				
	PLACE1011160	3.37	3.04	1.43	3.14	2.88	3.52	2.47	3.09	3.09				
	PLACE1011165	2.82	1.49	1.92	2.09	2.17	1.87	1.17	1.03	1.03				
	PLACE1011181	4.06	3.32	2.04	6.31	7.78	3.19	4.22	5.69	5.69				
	PLACE1011185	3.65	1.45	1.75	3.91	3.68	3.21	2.20	1.92	1.92				
	PLACE1011186	10.21	6.77	9.51	8.05	10.88	9.70	7.96	8.75	8.75				
50	PLACE1011203	0.72	0.41	0.60	0.86	1.05	1.05	0.76	0.91	0.91	*	+		
	PLACE1011214	2.12	1.28	1.87	3.46	4.03	3.52	3.00	2.98	2.98	**	+	**	+
	PLACE1011219	5.09	4.32	4.64	6.07	4.80	4.51	3.57	4.01	4.01		*	-	
	PLACE1011221	8.97	5.27	6.20	7.22	3.06	12.50	3.65	4.01	4.01				
	PLACE1011229	3.75	1.90	2.65	1.71	3.17	2.00	1.40	2.18	2.18				
55	PLACE1011231	3.92	2.25	2.28	3.91	4.79	2.77	5.22	2.91	2.91				
	PLACE1011236	8.67	4.88	5.11	4.86	4.69	4.64	5.23	5.25	5.25				
	PLACE1011247	4.61	2.86	4.14	4.95	3.62	3.74	4.69	6.05	6.05				

Table 327

5	PLACE1010410	4.61	1.87	2.21	5.83	8.60	7.04	3.70	4.91	4.91	*	+		
	PLACE1010418	3.29	1.76	2.41	6.21	6.34	6.33	2.79	3.38	3.38	**	+		
	PLACE1010425	1.18	0.35	0.46	1.22	0.78	1.70	0.80	1.37	1.37				
	PLACE1010443	5.43	3.03	3.71	5.62	3.76	6.48	4.57	5.05	5.05				
	PLACE1010445	4.33	2.64	3.67	5.95	5.86	6.97	4.11	3.20	3.2	*	+		
10	PLACE1010481	1.37	1.21	1.06	0.8	0.77	1.60	1.13	1.10	1.1				
	PLACE1010482	5.16	2.61	3.60	3.41	3.22	3.80	5.36	2.91	2.91				
	PLACE1010491	2.88	2.21	3.23	5.03	5.64	4.25	5.35	8.41	8.41	*	+	*	+
	PLACE1010492	2.47	1.94	1.90	1.59	2.93	3.57	2.66	2.46	2.46				
	PLACE1010509	1.31	0.33	0.65	0.44	1.02	0.95	1.07	1.11	1.11				
15	PLACE1010518	4.3	2.12	3.06	8.55	9.22	8.31	5.08	9.10	9.1	**	+	*	+
	PLACE1010522	4.42	3.30	2.99	4.43	3.15	5.70	4.02	5.51	5.51				
	PLACE1010529	4.44	3.27	3.34	4.15	2.17	4.43	2.83	4.60	4.6				
	PLACE1010547	1.36	0.46	1.84	1.38	2.57	0.83	0.81	0.68	0.68				
	PLACE1010560	3.62	1.42	1.78	3.44	4.11	3.17	1.69	3.25	3.25				
	PLACE1010562	2.49	1.56	1.51	2.33	1.85	1.73	1.62	1.70	1.7				
20	PLACE1010579	1.43	1.21	2.19	1.9	1.92	3.18	1.68	1.93	1.93				
	PLACE1010590	6.35	2.50	3.66	4.91	4.74	4.81	3.94	5.30	5.3				
	PLACE1010599	2.99	2.56	2.79	4.69	2.68	4.02	2.68	2.87	2.87				
	PLACE1010606	0.64	1.41	0.70	0.91	1.32	1.04	0.85	0.75	0.75				
	PLACE1010616	1.07	0.75	1.12	3.22	1.83	3.57	1.94	1.36	1.36	*	+		
25	PLACE1010622	9.24	4.26	4.31	2.37	3.79	2.39	1.80	2.04	2.04				
	PLACE1010624	6.73	4.32	4.19	2.38	2.68	1.71	1.73	1.83	1.83	*	-	*	-
	PLACE1010628	1.26	1.28	1.00	1.32	2.31	1.21	1.10	0.98	0.98				
	PLACE1010629	1.86	1.74	1.96	1.86	4.02	4.33	2.68	2.28	2.28		*	+	
	PLACE1010630	5.11	3.33	3.71	7.92	7.09	5.39	5.90	7.29	7.29	*	+	*	+
30	PLACE1010631	1.79	0.95	0.97	2.41	2.47	2.83	1.91	1.86	1.86	*	+		
	PLACE1010651	2.68	2.44	2.01	2.53	1.74	2.28	2.68	4.49	4.49				
	PLACE1010661	2.42	1.52	2.69	2.28	2.26	4.08	1.65	3.04	3.04				
	PLACE1010662	2.49	1.93	2.59	3.46	2.35	2.86	1.94	1.49	1.49				
	PLACE1010668	6.55	2.72	2.43	7.07	8.23	6.07	5.21	6.36	6.36				
	PLACE1010702	18.26	8.81	10.62	33.41	42.20	27.93	11.82	16.20	16.2	*	+		
35	PLACE1010709	29.25	14.24	17.35	21.38	21.56	17.73	31.21	41.95	41.95		*	+	
	PLACE1010713	11.16	4.98	5.23	9.06	10.03	9.81	9.13	15.19	15.19				
	PLACE1010714	0.55	0.48	0.52	0.64	0.75	1.34	0.77	0.70	0.7		**	+	
	PLACE1010716	5.99	2.36	2.79	3.78	2.95	5.02	3.07	3.15	3.15				
	PLACE1010717	2.06	1.35	1.59	2.22	1.80	2.83	0.90	1.52	1.52				
40	PLACE1010720	18.67	8.95	8.08	12.05	17.26	10.51	4.13	4.57	4.57				
	PLACE1010739	1.36	1.32	0.50	2.03	2.27	3.00	2.05	1.94	1.94	*	+	*	+
	PLACE1010743	1.84	1.21	0.69	1.5	0.87	0.37	0.87	1.50	1.5				
	PLACE1010752	5.21	2.95	2.72	2.98	3.08	1.69	2.31	3.98	3.98				
	PLACE1010761	9.42	7.63	8.64	20.89	19.08	19.20	8.58	11.68	11.68	**	+		
45	PLACE1010771	7.47	3.15	3.53	5.95	5.91	7.07	6.15	6.64	6.64				
	PLACE1010784	0.87	0.52	1.39	0.62	1.10	1.01	1.14	0.89	0.89				
	PLACE1010786	3.62	2.60	1.59	2.95	1.86	4.15	2.62	2.64	2.64				
	PLACE1010789	2.47	1.71	1.29	7.34	5.69	4.59	3.94	2.83	2.83	**	+		
	PLACE1010800	5.09	2.34	2.77	6.42	5.52	4.71	4.26	4.86	4.86				
	PLACE1010802	2.85	0.65	1.48	2.19	2.46	1.85	2.00	3.34	3.34				
50	PLACE1010811	3.15	1.56	1.75	2.32	2.73	2.52	1.52	3.78	3.78				
	PLACE1010813	4.37	2.54	2.23	3.08	2.72	2.68	2.51	3.14	3.14				
	PLACE1010827	2.09	0.81	0.76	1.38	1.83	1.70	1.14	4.49	4.49				
	PLACE1010833	6.2	2.64	3.19	13.01	9.81	9.59	5.31	5.99	5.99	*	+		
	PLACE1010839	3.43	2.31	3.38	7.65	5.31	7.09	3.40	4.37	4.37	*	+		
	PLACE1010856	3.15	2.01	1.95	2.4	2.08	1.61	2.50	2.16	2.16				
55	PLACE1010857	5.31	2.37	3.64	2.66	4.62	2.73	2.23	4.31	4.31				
	PLACE1010870	6.19	2.76	3.14	7.02	8.56	7.22	4.56	4.25	4.25	*	+		

Table 326

5	PLACE1010045	6.37	3.18	4.06	5.76	9.98	5.99	3.46	7.87	7.87			
	PLACE1010053	7.31	4.10	4.89	8.33	10.67	7.68	5.23	4.89	4.89			
	PLACE1010060	5.81	2.55	2.85	4.53	3.83	3.76	4.23	4.25	4.25			
	PLACE1010069	1.38	1.53	1.33	0.77	1.42	0.61	0.88	2.96	2.96			
	PLACE1010070	1.16	0.11	0.64	0.75	0.45	1.16	2.70	1.27	1.27			
	PLACE1010074	9.55	3.59	4.51	8.29	7.15	7.46	5.88	9.16	9.16			
10	PLACE1010076	32.02	14.06	13.18	16.2	20.29	12.88	25.05	26.03	26.03			
	PLACE1010078	5.69	2.44	3.34	4.22	3.97	3.39	4.99	6.24	6.24			
	PLACE1010081	3.3	1.78	4.36	4.28	4.59	3.29	2.67	5.51	5.51			
	PLACE1010083	2.72	1.96	1.66	0.92	1.44	1.20	2.07	2.63	2.63			
	PLACE1010089	2.82	1.29	2.28	4.53	3.47	5.64	3.30	4.44	4.44	*	+	+
15	PLACE1010096	3.39	1.17	2.00	2.56	2.19	2.70	1.45	1.92	1.92			
	PLACE1010102	5.26	3.31	3.97	9.27	6.87	8.63	4.86	8.37	8.37	*	+	
	PLACE1010105	4.29	0.95	1.09	2.44	2.73	1.94	2.71	4.01	4.01			
	PLACE1010106	1.98	0.97	0.87	3.59	2.61	2.19	3.70	3.99	3.99	*	+	**
	PLACE1010130	2.14	1.13	1.35	4.01	3.52	4.49	5.26	8.14	8.14	**	+	**
20	PLACE1010132	6.25	4.26	5.07	4.52	4.25	5.01	4.63	5.39	5.39			
	PLACE1010134	3.87	1.25	2.18	2.61	2.68	1.90	2.26	2.82	2.82			
	PLACE1010139	28.44	17.86	18.97	14.51	12.81	13.26	30.01	30.01	30.01			
	PLACE1010148	2.71	1.27	1.28	1.81	1.73	1.69	1.33	1.07	1.07			
	PLACE1010152	2.7	1.53	1.95	4.96	4.00	5.90	3.04	3.45	3.45	*	+	*
25	PLACE1010155	1.95	0.77	1.06	1.99	1.84	1.65	3.04	2.97	2.97		**	+
	PLACE1010156	1.86	1.01	1.72	5.69	7.58	4.30	7.96	8.94	8.94	*	+	**
	PLACE1010161	2.56	0.74	1.26	2.69	3.12	1.69	2.27	2.44	2.44			
	PLACE1010181	1.28	0.65	2.02	2	2.26	1.95	1.65	3.46	3.46			
	PLACE1010194	4.75	3.52	3.08	5.35	3.77	3.54	4.56	3.30	3.3			
	PLACE1010202	1.47	0.70	0.65	1.46	1.47	1.29	1.34	1.69	1.69			
30	PLACE1010231	1.3	1.19	0.99	2.11	1.60	1.20	1.89	1.43	1.43			
	PLACE1010235	2.55	0.79	1.71	2.65	2.39	3.67	1.07	1.49	1.49			
	PLACE1010237	0.84	1.17	0.50	1.96	1.56	2.36	1.09	0.99	0.99	*	+	
	PLACE1010251	3.81	2.13	2.41	3.72	3.24	1.88	1.45	3.83	3.83			
	PLACE1010261	1.35	0.55	0.65	1.04	1.71	1.55	1.14	1.11	1.11			
35	PLACE1010270	1.46	0.23	0.71	1.47	1.36	1.19	1.45	1.50	1.5			
	PLACE1010273	0.99	0.27	0.37	1.03	1.00	0.75	1.88	1.40	1.4		*	+
	PLACE1010274	5.85	2.65	3.07	9.77	6.41	6.98	9.03	7.48	7.48		*	+
	PLACE1010277	0.73	0.48	1.84	2.72	1.75	2.20	2.90	4.07	4.07		**	+
	PLACE1010293	2.98	2.04	1.13	2.91	3.54	3.25	2.77	2.03	2.03			
	PLACE1010297	1.4	1.02	0.95	3.02	1.83	2.84	1.39	2.38	2.38	*	+	
40	PLACE1010300	2.53	1.14	1.11	3.81	3.04	2.55	5.33	3.77	3.77		*	+
	PLACE1010310	32.51	17.93	15.91	30.53	26.14	27.60	23.13	27.43	27.43			
	PLACE1010321	4.23	1.98	2.58	2.3	3.07	2.72	3.25	3.30	3.3			
	PLACE1010324	1.39	0.54	0.66	1.12	1.26	0.93	0.53	1.22	1.22			
	PLACE1010329	2.31	0.98	1.09	3.01	2.16	2.51	0.92	2.83	2.83			
45	PLACE1010330	5.03	4.25	4.39	4.99	4.21	5.53	4.14	7.36	7.36			
	PLACE1010335	15.88	12.79	14.20	8.65	7.50	7.75	5.10	7.02	7.02	**	-	**
	PLACE1010341	0.29	0.66	0.37	0.99	1.99	0.90	0.42	0.24	0.24			
	PLACE1010342	0.95	0.44	0.79	1.64	1.38	0.95	1.00	0.59	0.59			
	PLACE1010346	4.09	1.92	1.71	4.43	3.58	3.75	2.92	3.05	3.05			
50	PLACE1010362	6.71	3.65	3.11	5.41	4.43	4.32	3.93	3.09	3.09			
	PLACE1010364	2.59	1.60	1.20	2.85	2.37	2.17	1.17	1.29	1.29			
	PLACE1010368	4.89	5.80	6.59	7.12	8.61	9.56	7.27	7.36	7.36	*	+	*
	PLACE1010373	5.27	3.94	4.28	6.3	5.08	7.04	5.11	6.77	6.77			
	PLACE1010383	4.96	2.31	2.06	8.93	6.77	7.08	4.88	5.39	5.39	*	+	
55	PLACE1010385	0.33	0.37	0.29	0.63	0.70	0.56	0.54	0.63	0.63	**	+	**
	PLACE1010389	5.32	1.69	2.00	3.34	2.68	4.99	2.13	3.58	3.58			
	PLACE1010401	1.04	0.68	0.65	0.51	0.65	1.60	0.87	0.66	0.66			

Table 325

5	PLACE1009607	3.67	1.38	1.49	4.1	6.22	4.84	3.49	3.02	3.02	*	+		
	PLACE1009613	3.3	1.40	2.05	3.5	4.21	3.44	2.36	3.25	3.25				
	PLACE1009621	2.39	1.87	2.42	5.45	5.01	5.43	4.10	6.00	6	**	+	**	+
	PLACE1009622	1.78	0.78	1.73	2.06	1.60	1.99	2.28	4.60	4.6			*	+
	PLACE1009624	0.78	1.54	0.90	2.28	3.24	1.75	2.30	1.54	1.54				
	PLACE1009637	1.33	0.77	0.84	3.69	2.89	3.73	3.44	3.33	3.33	**	+	**	+
10	PLACE1009639	2.08	0.08	0.65	2.19	1.89	1.62	1.82	1.57	1.57				
	PLACE1009654	2.53	0.76	1.11	2.83	1.57	1.39	1.86	2.31	2.31				
	PLACE1009659	5.89	3.14	3.71	3.85	5.36	4.32	4.03	5.80	5.8				
	PLACE1009665	1.27	1.04	0.92	2.92	2.14	2.54	0.79	2.03	2.03	**	+		
	PLACE1009669	3.5	3.60	3.11	3.69	5.37	3.54	3.97	4.99	4.99			*	+
15	PLACE1009670	2.16	1.80	1.32	3.29	1.88	3.37	1.92	2.64	2.64				
	PLACE1009708	2.48	1.90	1.93	4.13	3.31	5.20	2.14	3.90	3.9	*	+		
	PLACE1009721	3.15	2.27	2.41	1.67	3.17	2.28	7.20	2.48	2.48				
	PLACE1009731	3.26	1.56	1.59	2.49	3.83	1.81	1.89	2.37	2.37				
	PLACE1009735	2.96	1.31	2.04	2.52	2.63	2.49	2.46	2.74	2.74				
20	PLACE1009737	2.94	0.82	1.29	2.21	2.29	2.41	1.51	1.54	1.54				
	PLACE1009741	3.13	1.21	2.06	2.99	2.40	4.38	1.51	3.07	3.07				
	PLACE1009752	3.23	1.55	1.75	2.3	2.72	2.29	1.86	1.61	1.61				
	PLACE1009763	5.82	2.68	2.79	4.62	5.11	4.63	5.66	4.98	4.98				
	PLACE1009766	1.66	0.72	1.60	4.14	2.26	2.27	1.82	1.34	1.34				
	PLACE1009772	1.8	1.13	2.05	2.49	1.48	2.20	2.00	2.91	2.91				
25	PLACE1009782	3.79	1.21	0.99	3.99	3.99	2.22	2.25	2.39	2.39				
	PLACE1009794	3.98	1.98	2.41	2.73	2.16	1.89	2.44	4.87	4.87				
	PLACE1009798	3.03	1.31	2.50	3.63	5.60	4.46	2.46	3.00	3	*	+		
	PLACE1009845	0.71	0.31	1.69	2.44	1.45	2.19	0.63	2.13	2.13				
	PLACE1009849	2.59	1.40	2.09	2.06	1.75	1.55	1.88	1.44	1.44				
30	PLACE1009857	2.54	1.21	2.06	1.63	1.90	1.80	2.01	3.22	3.22				
	PLACE1009861	3.24	2.05	2.05	5.01	4.66	4.82	3.10	3.89	3.89	**	+		
	PLACE1009872	43.66	21.33	23.44	30.54	23.07	32.80	14.91	18.35	18.35				
	PLACE1009877	34.76	13.19	14.79	13.63	20.45	13.77	10.79	13.80	13.8				
	PLACE1009879	1.98	0.47	1.85	1.36	3.33	1.12	1.96	1.87	1.87				
35	PLACE1009886	1.09	0.42	0.92	1.49	1.32	1.87	0.94	1.34	1.34	*	+		
	PLACE1009888	3.11	1.53	2.24	1.6	2.71	2.32	1.87	2.30	2.3				
	PLACE1009908	4.53	2.06	2.64	3.65	2.87	3.85	3.36	4.12	4.12				
	PLACE1009919	5.7	2.20	3.89	5.91	4.05	5.41	4.60	6.30	6.3				
	PLACE1009921	1.24	0.74	1.00	0.94	2.00	1.75	1.08	0.94	0.94				
40	PLACE1009923	2.95	1.00	1.09	2.18	1.25	5.57	0.84	2.57	2.57				
	PLACE1009924	4.78	1.22	4.05	2.57	4.25	2.76	1.54	3.00	3				
	PLACE1009925	1.27	0.73	0.91	0.45	0.87	0.31	1.52	2.61	2.61			*	+
	PLACE1009931	11.44	4.02	5.58	10.31	11.46	9.16	5.01	7.71	7.71				
	PLACE1009935	0.24	0.55	0.45	0.68	0.48	0.50	1.11	1.18	1.18			**	+
	PLACE1009947	4.92	1.59	1.73	2.29	3.03	2.70	3.05	3.68	3.68				
45	PLACE1009961	1.11	1.73	1.45	1.96	1.63	2.02	1.58	0.96	0.96				
	PLACE1009971	2.28	1.16	1.31	3.83	2.51	3.34	2.27	2.89	2.89	*	+		
	PLACE1009982	7.21	2.79	4.22	5.07	5.60	7.20	6.47	7.74	7.74				
	PLACE1009992	3.36	1.01	0.95	2.29	3.14	1.23	2.29	4.00	4				
	PLACE1009995	7.97	4.77	4.17	10.64	14.64	12.97	7.62	12.10	12.1	*	+		
50	PLACE1009997	3.62	1.37	1.19	4.05	4.29	3.62	2.02	2.74	2.74				
	PLACE1010002	3.23	0.90	2.15	1.8	3.62	1.37	1.28	2.45	2.45				
	PLACE1010011	3.01	1.89	1.75	1.26	1.49	1.10	1.92	1.85	1.85				
	PLACE1010013	1.67	0.86	0.88	1.15	1.55	0.74	1.18	1.56	1.56				
	PLACE1010021	2.43	0.87	2.19	2.61	2.46	2.89	3.71	2.58	2.58				
55	PLACE1010023	4.84	1.80	2.28	2.57	2.67	3.95	1.95	4.34	4.34				
	PLACE1010031	5.58	2.99	1.54	4.23	4.90	2.66	3.18	2.93	2.93				
	PLACE1010039	1.86	0.50	0.58	0.41	1.70	1.28	1.06	1.35	1.35				

Table 324

5	PLACE1009190	2.12	1.27	2.18	1.35	2.00	2.47	0.78	2.21	2.21				
	PLACE1009196	1.64	0.69	1.48	2.04	2.57	3.98	1.85	1.52	1.52				
	PLACE1009200	4.32	1.99	2.61	4.48	5.35	4.97	2.74	2.68	2.68				
	PLACE1009217	2.54	0.82	0.83	0.92	1.24	1.76	2.27	2.78	2.78				
	PLACE1009230	3.29	1.25	2.57	3.85	3.86	4.23	1.77	4.02	4.02				
	PLACE1009236	3.68	1.44	1.56	2.57	2.82	2.63	1.54	2.09	2.09				
10	PLACE1009246	9.73	3.62	4.17	6.98	7.72	5.06	6.33	5.96	5.96				
	PLACE1009245	21.04	8.85	7.61	12.85	14.86	12.34	4.96	7.60	7.6				
	PLACE1009279	1.84	0.86	0.79	1.58	1.52	1.53	1.15	1.01	1.01				
	PLACE1009298	3.7	2.72	2.61	7.54	8.77	8.06	7.00	9.82	9.82	**	+	**	+
	PLACE1009308	8.08	4.61	4.25	6.42	4.02	4.44	5.48	7.05	7.05				
15	PLACE1009319	2.03	1.05	1.47	2.87	1.77	3.10	1.90	2.70	2.7				
	PLACE1009328	1.59	0.99	1.42	4.54	4.75	5.66	3.66	4.23	4.23	**	+	**	+
	PLACE1009335	1.22	0.54	0.61	2.18	1.74	1.92	1.46	0.54	0.54	**	+		
	PLACE1009338	3.48	1.35	1.84	5.85	6.71	4.36	2.31	2.98	2.98	*	+		
	PLACE1009344	3.01	1.13	2.79	1.83	3.29	2.00	2.97	2.70	2.7				
20	PLACE1009355	1.86	0.75	0.42	1.64	1.55	1.14	2.65	5.34	5.34		*	+	
	PLACE1009368	2.14	1.43	1.26	1.31	1.41	1.74	1.22	2.07	2.07				
	PLACE1009375	1.44	0.73	1.31	0.98	2.28	1.80	1.47	2.25	2.25				
	PLACE1009388	1.69	1.27	1.19	3.96	2.82	3.05	1.65	2.75	2.75	**	+		
	PLACE1009398	6.96	2.57	3.77	9	5.66	6.33	4.19	4.18	4.18				
25	PLACE1009404	4.11	2.25	3.40	3.14	5.18	4.09	2.94	3.62	3.62				
	PLACE1009410	1.58	0.66	0.54	0.77	1.47	0.75	1.04	1.03	1.03				
	PLACE1009417	1.85	0.80	1.11	2.36	1.87	0.83	1.31	3.04	3.04				
	PLACE1009424	10.71	5.65	7.84	8.47	7.50	6.48	8.06	10.17	10.17				
	PLACE1009434	3.29	1.53	1.47	2.38	1.85	1.49	1.58	1.71	1.71				
30	PLACE1009443	2.96	1.10	1.13	1.36	1.62	1.85	0.98	1.60	1.6				
	PLACE1009444	3.55	2.71	1.84	4.89	4.13	5.32	3.26	4.47	4.47	*	+		
	PLACE1009459	5.23	2.29	2.82	3.92	3.20	3.43	3.08	4.21	4.21				
	PLACE1009460	0.43	0.37	0.33	0.44	0.37	1.88	0.42	0.69	0.69				
	PLACE1009468	5.92	2.35	2.32	5.44	2.65	2.84	4.15	2.97	2.97				
	PLACE1009476	2.6	0.89	1.54	2.02	2.17	1.83	1.69	2.92	2.92				
35	PLACE1009477	3.84	1.44	1.65	4.37	2.97	3.00	2.09	2.93	2.93				
	PLACE1009493	2.08	0.70	1.33	2.12	1.22	1.24	0.82	2.09	2.09				
	PLACE1009502	0.95	0.44	0.76	0.93	0.90	0.72	0.86	1.97	1.97				
	PLACE1009524	2.21	0.79	1.36	1.49	1.81	1.15	1.58	2.15	2.15				
	PLACE1009527	1.81	1.71	1.43	2.21	1.72	1.29	1.43	1.91	1.91				
40	PLACE1009531	5.24	3.01	2.51	5.69	3.69	5.37	6.78	6.24	6.24		*	+	
	PLACE1009535	1.5	0.44	0.55	2.44	1.80	1.98	2.38	1.44	1.44	*	+		
	PLACE1009539	3.39	1.25	2.38	2.92	3.40	3.47	2.40	3.54	3.54				
	PLACE1009540	6	3.37	5.39	4.83	4.41	4.48	4.99	6.08	6.08				
	PLACE1009542	2.35	1.42	1.51	1.82	1.71	1.38	1.98	2.97	2.97				
45	PLACE1009546	1.47	0.53	0.69	0.94	1.26	0.62	1.78	0.85	0.85				
	PLACE1009556	1.35	0.95	1.07	1.35	2.21	0.98	2.07	2.20	2.2		**	+	
	PLACE1009569	2.13	1.30	1.80	2.87	2.82	3.16	1.76	2.07	2.07	*	+		
	PLACE1009571	2.72	1.88	1.50	2.08	1.82	2.50	1.30	1.73	1.73				
	PLACE1009573	8.32	4.58	4.70	4.98	4.59	3.46	3.53	2.68	2.68				
	PLACE1009576	3.44	1.43	2.32	5.25	5.67	4.91	3.85	4.08	4.08	**	+		
50	PLACE1009580	2.8	1.13	1.78	3.7	3.05	2.82	3.23	3.94	3.94		*	+	
	PLACE1009581	2.06	1.05	0.67	2.39	2.40	2.16	4.59	3.83	3.83		**	+	
	PLACE1009587	1.75	1.08	0.69	1.11	1.37	1.41	1.48	2.01	2.01				
	PLACE1009593	2.92	1.61	2.66	2.04	2.95	2.35	2.20	2.52	2.52				
	PLACE1009595	4.18	2.88	2.05	6.18	7.63	5.15	3.71	4.73	4.73	*	+		
55	PLACE1009596	1.65	1.09	0.96	1.87	2.16	2.45	1.64	1.48	1.48	*	+		
	PLACE1009600	6.27	3.95	2.87	7.95	6.45	4.31	4.05	5.38	5.38				
	PLACE1009604	2.52	0.69	0.99	3.24	2.45	1.67	2.85	2.11	2.11				

Table 323

5	PLACE1008738	1.83	2.28	2.00	1.8	1.24	1.00	1.36	3.09	3.09				
	PLACE1008742	4.02	1.70	1.54	4.3	5.17	3.46	2.80	3.04	3.04				
	PLACE1008744	1.17	0.49	0.67	1.04	1.21	1.19	1.03	1.69	1.69				
	PLACE1008748	1.18	0.53	1.02	1.35	1.38	1.66	1.55	1.10	1.1				
	PLACE1008757	0.57	0.66	1.64	0.96	1.31	1.19	0.28	1.35	1.35				
10	PLACE1008766	5.2	1.84	3.38	5.73	6.06	11.79	4.24	3.09	3.09				
	PLACE1008785	3.43	1.55	1.67	3.73	3.48	3.51	2.86	2.40	2.4				
	PLACE1008790	4.68	2.15	2.15	5.43	4.49	3.61	3.28	3.45	3.45				
	PLACE1008798	6.35	0.62	2.86	2.36	3.47	2.89	1.71	2.65	2.65				
	PLACE1008807	0.99	1.20	1.36	0.98	1.48	1.58	0.90	2.29	2.29				
15	PLACE1008808	2.02	1.19	1.16	1.26	1.76	1.00	2.24	1.72	1.72				
	PLACE1008813	0.94	0.76	1.96	0.73	1.40	0.71	0.81	2.94	2.94				
	PLACE1008836	3.35	2.03	2.82	3.36	3.83	3.93	1.76	4.97	4.97				
	PLACE1008851	6.7	2.37	2.20	3.21	3.73	4.45	1.84	2.02	2.02				
	PLACE1008854	1.01	0.67	0.67	0.73	1.08	1.01	0.89	0.70	0.7				
	PLACE1008864	5.23	2.45	2.26	6.92	5.09	5.19	3.11	3.68	3.68				
20	PLACE1008867	1.96	1.55	1.26	5.74	4.65	5.92	4.30	4.51	4.51	**	+	**	+
	PLACE1008876	51.43	26.54	27.05	38	43.35	42.72	24.30	22.52	22.52				
	PLACE1008887	1.78	0.54	1.07	2.31	2.39	2.93	1.78	2.61	2.61	*	+		
	PLACE1008902	1.97	0.82	0.85	1.66	1.42	3.56	1.02	2.90	2.9				
	PLACE1008911	6.01	5.11	5.63	8.6	8.99	8.79	6.07	6.33	6.33	**	+		
25	PLACE1008917	3.34	2.37	2.25	2.83	3.74	3.27	2.99	3.43	3.43				
	PLACE1008920	1.37	0.52	0.53	1.3	2.33	1.36	0.77	1.37	1.37				
	PLACE1008925	1.43	1.01	0.48	2.16	1.60	0.85	1.24	0.93	0.93				
	PLACE1008930	8.48	4.04	4.74	5.59	5.27	6.20	2.97	5.51	5.51				
	PLACE1008934	2.73	1.83	1.68	2.96	2.07	1.68	2.13	1.92	1.92				
30	PLACE1008941	2.12	2.49	2.29	2.81	3.70	3.18	1.74	1.69	1.69	*	+	**	-
	PLACE1008947	5.3	4.86	3.97	6.01	5.96	5.46	4.91	5.47	5.47				
	PLACE1008984	2.32	1.08	1.90	4.47	4.44	4.99	1.56	2.13	2.13	**	+		
	PLACE1008985	1.06	1.41	1.57	2.31	2.24	1.90	1.29	3.49	3.49	*	+		
	PLACE1008994	1.26	0.32	0.61	1.19	2.34	0.75	0.51	0.61	0.61				
	PLACE1009020	2.03	0.83	0.79	1.36	0.98	0.99	0.91	1.17	1.17				
35	PLACE1009027	2.42	0.29	0.98	17.03	20.58	24.13	13.27	17.48	17.48	**	+	**	+
	PLACE1009039	0.66	0.39	0.60	0.97	0.77	0.82	0.81	1.68	1.68	*	+	*	+
	PLACE1009045	1.25	0.20	1.18	0.92	1.61	1.30	3.10	3.19	3.19			**	+
	PLACE1009048	0.29	0.37	0.55	0.51	0.66	0.96	1.13	0.67	0.67				
	PLACE1009050	0.48	(0.04)	0.53	1.13	0.72	1.09	0.42	0.86	0.86	*	+		
40	PLACE1009060	3.31	1.27	1.72	4.36	1.92	4.74	2.50	4.91	4.91				
	PLACE1009067	4.9	1.27	1.78	2.92	1.97	2.26	4.68	4.77	4.77				
	PLACE1009071	5.93	4.97	3.58	6.84	4.81	6.47	5.46	4.55	4.55				
	PLACE1009090	3.14	0.90	2.12	3.01	2.91	5.24	2.46	1.95	1.95				
	PLACE1009091	4.11	1.05	1.26	1.69	2.73	1.26	0.58	1.98	1.98				
45	PLACE1009094	2.34	2.30	1.26	2.48	1.83	1.50	3.22	2.13	2.13				
	PLACE1009099	4.71	2.33	2.35	5.94	5.89	8.61	3.69	5.79	5.79	*	+		
	PLACE1009110	1.06	1.13	0.63	4.86	1.21	3.08	2.60	2.41	2.41		**	+	
	PLACE1009111	1.61	0.55	0.64	2.6	1.26	1.76	1.01	2.06	2.06				
	PLACE1009113	5.16	1.93	2.40	3.84	4.61	2.47	1.71	4.56	4.56				
	PLACE1009130	2.4	1.03	1.11	1.45	1.93	2.94	1.60	1.65	1.65				
50	PLACE1009150	1.73	0.66	1.55	2.16	2.01	2.30	1.65	1.47	1.47				
	PLACE1009155	3.13	2.31	1.89	4.69	5.44	5.47	2.82	2.95	2.95	**	+		
	PLACE1009158	3.54	1.36	1.91	2.88	2.25	2.53	2.92	2.13	2.13				
	PLACE1009166	2.58	1.73	2.09	2.03	2.15	2.17	2.50	2.39	2.39				
	PLACE1009172	2.84	0.78	1.90	4.25	3.46	3.67	2.50	4.14	4.14	*	+		
55	PLACE1009174	3.1	1.74	1.40	4.47	5.90	4.15	2.46	2.47	2.47	*	+		
	PLACE1009183	6.02	1.51	2.01	3.8	4.98	3.03	2.18	2.53	2.53				
	PLACE1009186	3.59	0.98	1.37	2.08	2.13	0.57	1.69	3.99	3.99				

Table 322

	PLACE1008330	3.99	1.02	3.12	3.69	2.72	3.55	2.59	3.30	3.3				
5	PLACE1008331	3.5	1.58	2.61	2.43	4.87	4.55	2.21	5.77	5.77				
	PLACE1008351	3.59	1.91	2.57	5.18	5.19	5.56	3.81	3.50	3.5	**	+		
	PLACE1008356	3.92	0.69	2.72	2.64	2.56	2.29	2.42	2.95	2.95				
	PLACE1008359	1.48	0.76	0.90	2.22	1.26	2.34	1.68	2.46	2.46		*	+	
10	PLACE1008368	4.18	1.66	2.15	9.15	7.54	8.92	6.11	7.44	7.44	**	+	**	+
	PLACE1008369	2.77	0.73	1.19	2.41	7.30	3.35	1.02	1.60	1.6				
	PLACE1008392	2.13	0.98	1.09	1.58	3.18	1.77	1.88	2.10	2.1				
	PLACE1008394	26.4	13.24	13.94	17.36	15.53	22.06	16.70	19.87	19.87				
	PLACE1008398	7.2	3.44	10.45	4.58	8.83	4.91	2.86	4.01	4.01				
	PLACE1008401	3.08	0.75	1.07	1.76	1.56	2.79	1.84	3.10	3.1				
15	PLACE1008402	6.01	1.01	4.48	2.49	3.09	3.48	2.05	3.35	3.35				
	PLACE1008405	25.84	13.96	18.38	38.51	28.28	49.12	27.91	33.39	33.39		*	+	
	PLACE1008409	16.67	9.55	11.29	12.69	10.07	15.56	12.51	11.76	11.76				
	PLACE1008420	5.7	4.00	2.86	5.32	4.44	3.71	4.42	4.23	4.23				
	PLACE1008424	3.57	2.25	1.23	2.09	2.46	2.00	2.48	2.25	2.25				
20	PLACE1008426	4.1	1.19	2.55	2.53	2.76	1.73	1.42	1.69	1.69				
	PLACE1008429	1.34	0.85	1.46	2	3.50	1.65	1.93	1.52	1.52				
	PLACE1008430	1.82	0.58	0.88	2.02	1.64	0.56	0.86	2.26	2.26				
	PLACE1008437	2.06	0.49	1.54	1.53	1.27	1.54	1.33	2.88	2.88				
	PLACE1008453	3.99	2.14	2.45	2.78	2.86	2.41	2.29	5.19	5.19				
	PLACE1008454	4.67	3.03	4.69	8.04	6.50	8.39	3.85	5.65	5.65	*	+		
25	PLACE1008455	6.35	2.17	1.87	10.14	10.23	5.77	6.05	5.82	5.82				
	PLACE1008457	9.43	3.52	3.32	5.83	7.73	6.63	5.24	7.01	7.01				
	PLACE1008465	2.14	1.13	1.61	1.55	3.02	1.33	2.20	2.70	2.7				
	PLACE1008469	12.37	7.23	7.87	8.96	9.09	12.38	13.17	10.93	10.93				
	PLACE1008488	1.94	0.92	1.25	0.9	1.06	1.44	1.44	0.95	0.95				
30	PLACE1008519	3.83	1.77	1.73	2.4	1.77	1.88	2.77	1.49	1.49				
	PLACE1008524	3.06	0.85	1.87	3.33	2.40	3.53	2.10	1.92	1.92				
	PLACE1008531	3.02	1.05	2.48	2.83	2.67	2.71	2.79	2.45	2.45				
	PLACE1008532	1.95	1.34	1.62	3.81	2.99	2.68	2.83	3.90	3.9	*	+	**	+
	PLACE1008533	6.08	2.16	3.15	4.18	5.64	3.25	3.67	5.24	5.24				
35	PLACE1008542	3.98	1.49	1.76	4.67	6.17	4.59	3.86	6.21	6.21	*	+		
	PLACE1008549	2.51	1.53	0.88	1.7	2.81	1.76	1.36	1.66	1.66				
	PLACE1008560	1.85	0.72	0.75	0.85	0.84	0.96	2.24	1.41	1.41				
	PLACE1008567	2.83	1.62	2.07	2.6	2.14	2.90	2.18	3.74	3.74				
	PLACE1008568	1.44	0.85	1.22	4.02	2.55	4.05	2.96	3.07	3.07	*	+	**	+
40	PLACE1008569	6.68	1.97	2.63	4.52	4.62	4.72	3.58	5.21	5.21				
	PLACE1008584	2.8	0.91	1.34	2.88	1.91	1.76	1.37	1.81	1.81				
	PLACE1008585	6.05	2.08	1.87	5.97	6.88	5.16	6.30	6.66	6.66				
	PLACE1008603	2.79	0.92	1.64	1.88	1.63	1.46	1.46	2.30	2.3				
	PLACE1008621	2.19	0.44	1.30	1.02	1.47	0.69	1.47	2.18	2.18				
45	PLACE1008625	0.9	0.37	0.80	0.8	0.79	0.63	1.51	1.36	1.36		*	+	
	PLACE1008626	1.01	0.36	0.40	1.03	0.59	0.80	0.60	2.30	2.3				
	PLACE1008627	3.31	1.35	1.85	3.04	2.64	2.27	2.82	2.83	2.83				
	PLACE1008629	4.46	2.86	3.88	4.95	3.86	3.87	2.66	4.45	4.45				
	PLACE1008630	6.49	3.28	4.20	4.75	4.80	4.92	3.62	3.61	3.61				
	PLACE1008643	3.94	1.90	2.23	4.63	3.91	2.95	3.01	3.94	3.94				
50	PLACE1008650	1.04	0.28	0.89	1.14	0.65	0.67	0.98	2.48	2.48				
	PLACE1008657	2.91	1.23	0.78	2.05	1.78	1.50	2.02	2.54	2.54				
	PLACE1008664	2.55	1.44	2.26	1.59	2.27	1.94	2.31	1.74	1.74				
	PLACE1008693	3.83	1.61	1.78	3.36	3.20	3.43	2.03	2.63	2.63				
	PLACE1008696	1.57	0.88	1.00	2.25	2.32	2.53	1.82	1.87	1.87	**	+	*	+
55	PLACE1008715	1.2	1.08	0.38	2.73	1.58	1.58	1.30	1.20	1.2				
	PLACE1008716	2.62	1.18	1.33	2.82	3.53	2.15	2.83	1.88	1.88				
	PLACE1008722	8.81	3.15	4.14	9.07	11.88	9.16	5.01	7.77	7.77				

Table 321

5	PLACE1007877	4.54	1.36	1.17	4.16	4.25	2.80	3.39	3.32	3.32				
	PLACE1007878	4.4	2.07	2.29	2.41	2.70	2.37	3.13	5.04	5.04				
	PLACE1007881	1.27	0.74	0.75	0.94	1.76	0.67	0.87	1.11	1.11				
	PLACE1007885	1.23	1.17	1.11	1.97	2.06	1.97	2.46	3.25	3.25	**	+	**	+
	PLACE1007897	2.56	0.68	1.11	1.75	1.79	1.50	1.00	2.88	2.88				
10	PLACE1007908	7.68	3.04	3.27	4.73	4.71	5.04	4.39	4.18	4.18				
	PLACE1007922	1.4	0.69	0.89	1.56	0.63	1.43	1.13	0.93	0.93				
	PLACE1007946	4.36	3.22	3.12	4.56	4.09	3.11	2.97	3.28	3.28				
	PLACE1007950	5.15	1.51	1.60	3.7	3.21	2.35	3.25	8.99	8.99				
	PLACE1007954	3.66	2.15	2.27	2.4	2.26	2.19	2.79	1.92	1.92				
	PLACE1007955	4.71	1.37	1.67	2.61	3.53	2.54	2.49	4.46	4.46				
15	PLACE1007956	4.42	1.04	2.64	3.61	3.50	3.32	2.21	3.84	3.84				
	PLACE1007958	1.93	0.27	1.12	1.34	1.94	1.66	1.60	1.84	1.84				
	PLACE1007965	2.55	1.76	1.99	2.32	2.51	3.02	1.19	2.52	2.52				
	PLACE1007969	6.03	2.86	2.43	4.73	5.79	6.79	4.72	3.77	3.77				
	PLACE1007971	3.53	1.27	2.02	3.82	4.31	3.71	3.34	3.31	3.31				
20	PLACE1007990	2.84	1.35	1.80	4.92	3.19	2.61	2.45	2.53	2.53				
	PLACE1008000	1.73	0.77	0.35	3.42	1.14	0.76	1.28	1.93	1.93				
	PLACE1008002	0.38	-0.09	0.23	1.64	0.83	0.73	1.52	1.90	1.9	*	+	**	+
	PLACE1008037	0.98	0.19	0.99	1.13	1.05	1.34	1.22	1.68	1.68				
	PLACE1008044	4.87	3.62	2.89	3.52	3.76	3.71	2.56	3.52	3.52				
25	PLACE1008045	1.81	1.03	1.31	1.51	1.59	1.22	1.49	2.12	2.12				
	PLACE1008080	4.1	3.05	2.36	3.11	3.91	2.99	2.39	3.89	3.89				
	PLACE1008092	2.02	1.71	1.46	1.1	0.88	0.81	1.07	2.15	2.15	*	-		
	PLACE1008095	2.93	1.27	1.19	2.55	1.83	2.32	1.34	3.34	3.34				
	PLACE1008105	2.48	0.98	1.47	2.27	0.97	1.49	2.91	5.54	5.54		*	+	
30	PLACE1008107	6.58	3.57	3.85	1.29	1.19	1.39	4.33	5.78	5.78	*	-		
	PLACE1008111	2.46	1.02	2.41	3.33	2.35	3.47	2.96	3.00	3				
	PLACE1008113	25.85	13.24	14.36	22.14	19.88	22.12	9.93	8.27	8.27				
	PLACE1008122	1.07	0.36	1.70	1.64	1.18	1.29	1.04	1.29	1.29				
	PLACE1008129	1.31	1.01	1.72	3.06	3.91	4.22	1.89	1.53	1.53	**	+		
	PLACE1008132	2.89	1.43	1.69	4.85	4.46	4.06	3.75	2.77	2.77	**	+		
35	PLACE1008137	3.98	1.85	1.77	2.91	2.34	1.96	2.43	2.78	2.78				
	PLACE1008174	10.37	5.11	6.06	7.46	7.08	5.83	3.58	4.68	4.68				
	PLACE1008177	5.22	2.35	2.42	4.78	5.45	4.55	2.08	2.73	2.73				
	PLACE1008181	0.6	0.35	0.59	2.1	1.63	0.83	0.78	0.73	0.73		*	+	
	PLACE1008195	4.21	3.69	4.41	3.34	3.31	4.29	3.54	5.03	5.03				
40	PLACE1008198	0.92	1.28	1.62	1.49	2.09	2.17	1.39	2.32	2.32				
	PLACE1008201	1.66	0.51	1.49	2.83	2.14	2.43	2.07	1.72	1.72	*	+		
	PLACE1008209	5.39	4.27	2.17	7.66	7.83	6.93	6.08	4.07	4.07	*	+		
	PLACE1008226	3.09	1.71	1.62	2.88	3.33	2.83	2.61	2.73	2.73				
	PLACE1008227	3.17	1.23	2.12	4.9	4.87	5.42	2.16	2.72	2.72	**	+		
45	PLACE1008231	2.12	0.50	0.70	1.87	1.47	1.28	1.21	0.99	0.99				
	PLACE1008238	3.15	3.76	3.38	3.65	4.20	4.83	4.89	4.62	4.62		**	+	
	PLACE1008244	1.2	0.39	0.55	1.23	1.76	1.25	0.99	1.37	1.37				
	PLACE1008249	2.18	0.27	0.99	2.07	1.35	1.47	0.79	1.16	1.16				
	PLACE1008266	3.92	3.58	3.56	7.01	10.06	9.34	6.60	6.76	6.76	**	+	**	+
	PLACE1008273	2.91	1.72	1.49	4.31	6.22	5.51	5.60	5.70	5.7	**	+	**	+
50	PLACE1008275	1.29	0.61	1.24	2.1	1.76	1.18	1.34	0.60	0.6				
	PLACE1008280	2.51	1.09	1.40	1.61	2.66	1.55	2.19	1.36	1.36				
	PLACE1008282	6.02	2.61	4.50	6.93	7.87	7.46	6.98	6.73	6.73	*	+		
	PLACE1008297	1.93	0.37	0.80	1.67	1.93	1.78	1.68	1.21	1.21				
	PLACE1008303	2.86	2.08	2.50	1.98	2.20	2.57	2.27	1.77	1.77				
55	PLACE1008309	1	0.36	0.94	1.24	0.57	1.29	1.65	0.87	0.87				
	PLACE1008315	12.99	6.08	6.62	5.63	6.15	4.10	5.15	5.08	5.08				
	PLACE1008329	5.4	1.64	1.66	3.46	3.09	2.21	1.61	3.12	3.12				

Table 320

	PLACE1007511	1.09	1.11	0.68	1.33	1.45	0.75	0.90	1.48	1.48				
5	PLACE1007513	4.69	1.71	2.94	3.5	3.66	3.78	3.32	6.37	6.37				
	PLACE1007524	6.92	2.48	2.90	3.93	4.08	2.82	1.80	1.66	1.66				
	PLACE1007525	4.99	2.20	2.97	4.48	5.31	5.23	2.35	2.30	2.3				
	PLACE1007537	3.67	3.75	2.72	3.67	3.58	4.70	2.62	4.19	4.19				
	PLACE1007544	1.23	1.96	1.26	3.11	3.23	2.88	3.01	2.55	2.55	**	+	*	+
10	PLACE1007547	3.83	2.63	2.50	6.49	5.11	5.77	2.96	2.23	2.23	**	+		
	PLACE1007557	3.78	2.86	3.01	6.18	5.42	6.26	3.20	3.81	3.81	**	+		
	PLACE1007560	7.5	4.33	3.69	5.21	4.40	3.63	6.61	8.29	8.29				
	PLACE1007565	1.39	0.57	0.51	1.55	0.69	1.08	1.27	0.93	0.93				
	PLACE1007580	0.78	0.25	0.56	1.38	0.71	0.94	1.33	1.46	1.46		**	+	
15	PLACE1007583	1.68	1.21	1.36	3.07	1.74	2.51	1.23	2.34	2.34				
	PLACE1007591	2.78	0.84	0.81	2.91	3.12	3.09	1.72	2.45	2.45				
	PLACE1007598	4.1	2.36	3.10	8.03	7.01	9.10	4.75	4.36	4.36	**	+		
	PLACE1007610	0.9	0.60	0.89	2.28	1.49	1.41	1.23	1.82	1.82	*	+	*	+
	PLACE1007618	1.76	1.24	1.15	1.76	2.07	1.52	1.03	1.29	1.29				
20	PLACE1007621	2.86	1.26	1.24	2.73	3.31	2.18	1.97	2.67	2.67				
	PLACE1007626	6.13	3.63	3.43	16.1	18.88	18.33	14.85	19.91	19.91	**	+	**	+
	PLACE1007632	4.92	2.23	3.27	3.4	3.01	3.01	4.94	4.29	4.29				
	PLACE1007635	3.04	0.96	2.65	2.16	2.56	2.69	1.76	2.94	2.94				
	PLACE1007645	4.04	1.20	2.15	4.72	5.27	5.01	4.78	4.87	4.87	*	+	*	+
25	PLACE1007649	1.28	0.79	0.67	1.29	1.36	2.38	1.28	2.15	2.15				
	PLACE1007659	4.23	1.93	2.69	6.75	3.97	6.88	2.94	4.41	4.41				
	PLACE1007669	6.2	1.80	2.99	5.47	6.53	4.51	3.57	2.86	2.86				
	PLACE1007677	4.22	1.89	1.71	6.84	8.75	7.28	3.90	4.46	4.46	**	+		
	PLACE1007688	5.22	1.69	2.55	2.63	3.33	2.71	2.38	2.43	2.43				
	PLACE1007690	3.97	2.16	3.39	4.09	4.66	3.97	3.53	4.50	4.5				
30	PLACE1007697	1.72	0.75	0.98	1.08	0.70	0.98	1.28	0.95	0.95				
	PLACE1007702	1.76	0.86	1.32	1.85	1.37	3.00	2.01	1.95	1.95				
	PLACE1007705	2.4	0.53	1.89	1.45	2.19	2.67	2.64	2.34	2.34				
	PLACE1007706	2.8	1.14	1.84	2.88	2.31	2.20	2.45	2.27	2.27				
	PLACE1007725	3.27	2.02	1.52	3.44	3.01	2.26	1.89	1.39	1.39				
35	PLACE1007729	3.75	0.91	0.48	1.28	1.88	1.09	1.35	1.46	1.46				
	PLACE1007730	4.12	1.63	2.33	3.92	2.43	2.55	1.94	4.18	4.18				
	PLACE1007737	4.58	2.53	1.58	4.31	5.53	6.14	3.60	3.45	3.45				
	PLACE1007743	1.47	0.73	0.61	2.7	2.78	2.53	1.94	2.71	2.71	**	+	*	+
	PLACE1007746	3.82	1.81	2.10	5.73	3.58	6.69	6.74	9.08	9.08		**	+	
40	PLACE1007753	2.19	1.29	1.71	1.02	1.20	1.89	1.49	1.55	1.55				
	PLACE1007769	0.98	0.53	0.69	1.58	1.14	1.77	1.01	1.04	1.04	*	+		
	PLACE1007780	4.5	2.26	1.99	3.89	4.09	2.46	2.36	2.20	2.2				
	PLACE1007791	5.12	2.18	2.04	3.75	4.60	3.26	2.31	3.66	3.66				
	PLACE1007807	2.35	0.20	1.17	3.74	3.71	3.65	3.45	3.14	3.14	*	+	*	+
45	PLACE1007810	1.24	0.07	0.47	1.06	0.82	1.32	1.17	1.10	1.1				
	PLACE1007814	5.26	2.80	2.95	4.73	4.47	4.22	4.12	5.14	5.14				
	PLACE1007828	1.64	1.27	1.04	1.35	1.67	1.95	1.48	2.89	2.89				
	PLACE1007829	6.87	2.06	4.61	11.59	10.29	14.12	4.54	6.24	6.24	*	+		
	PLACE1007841	2.09	0.69	0.83	1.22	2.33	3.41	1.28	2.06	2.06				
	PLACE1007842	2.47	1.09	2.35	2.63	2.75	2.08	1.49	2.36	2.36				
50	PLACE1007843	1.12	0.63	0.54	0.94	1.58	0.91	0.72	0.88	0.88				
	PLACE1007845	3.75	1.17	2.10	1.73	2.80	1.91	2.13	2.52	2.52				
	PLACE1007846	4.22	1.07	1.41	5.34	3.41	3.57	3.77	4.73	4.73				
	PLACE1007848	1.96	0.65	0.52	1.52	2.21	2.88	2.45	2.38	2.38		*	+	
	PLACE1007852	2.98	0.96	2.30	2.1	4.32	3.04	2.76	3.44	3.44				
55	PLACE1007858	1.43	0.68	1.60	6.59	6.03	5.72	4.91	4.91	4.91	**	+	**	+
	PLACE1007866	30.58	17.58	17.98	11.58	9.80	8.70	10.80	10.41	10.41				
	PLACE1007871	22.99	9.51	12.34	18.55	16.06	17.86	13.04	17.83	17.83				

Table 319

5	PLACE1007092	13.8	11.82	5.85	6.03	7.76	3.70	4.55	4.38	4.38				
	PLACE1007096	3.67	1.72	2.42	3.85	3.61	3.33	2.77	3.95	3.95				
	PLACE1007097	2.22	0.99	0.99	1.67	2.32	2.35	2.32	1.09	1.09				
	PLACE1007099	3.21	1.35	2.99	3.75	3.60	3.90	2.21	4.60	4.6				
	PLACE1007105	3.27	1.47	1.70	2.02	1.66	2.46	3.10	2.81	2.81				
10	PLACE1007108	1.84	0.54	0.64	1.21	1.32	0.77	1.03	1.13	1.13				
	PLACE1007111	1.12	0.75	0.77	2.41	0.87	1.64	1.17	1.43	1.43		*		+
	PLACE1007112	2.23	1.33	1.93	1.71	1.54	2.89	1.30	2.04	2.04				
	PLACE1007130	1.72	0.36	0.26	1	1.71	0.63	0.85	1.29	1.29				
	PLACE1007132	3.87	1.51	1.93	3.65	4.98	3.98	2.58	2.83	2.83				
15	PLACE1007140	2.78	1.67	1.49	5.51	4.02	1.95	1.59	4.61	4.61				
	PLACE1007143	4.57	2.06	2.35	3.69	3.88	3.45	2.67	3.35	3.35				
	PLACE1007169	7.86	3.91	6.07	4.6	3.97	4.34	4.66	5.06	5.06				
	PLACE1007178	3.63	1.78	2.11	3.46	2.58	2.44	3.58	4.50	4.5				
	PLACE1007190	1.52	0.85	1.18	1.02	0.96	1.35	1.62	1.51	1.51				
	PLACE1007201	1.85	0.34	1.11	1.37	0.91	2.07	0.93	1.05	1.05				
20	PLACE1007202	18.73	9.75	12.22	19.49	17.57	13.05	23.70	22.24	22.24		*		+
	PLACE1007226	4.6	2.18	1.44	3.72	3.17	3.32	4.10	4.25	4.25				
	PLACE1007238	4.59	1.78	4.87	4.05	4.43	2.63	3.54	2.85	2.85				
	PLACE1007239	4.19	2.58	2.67	5.05	3.84	2.86	3.07	4.50	4.5				
	PLACE1007242	3.6	1.20	1.84	1.27	2.10	2.41	1.99	2.58	2.58				
25	PLACE1007243	10.2	5.01	6.25	4.24	5.71	6.21	7.36	6.08	6.08				
	PLACE1007247	3.28	2.10	1.67	14.75	8.63	15.61	4.03	8.60	8.6	**	+	*	+
	PLACE1007257	7.61	5.72	7.16	3.66	3.64	3.79	1.96	3.64	3.64	**	-	**	-
	PLACE1007274	4.38	2.42	3.36	7.38	8.79	6.79	3.07	4.64	4.64	**	+		
	PLACE1007276	2.97	1.43	1.54	2.93	2.81	2.34	1.57	3.92	3.92				
30	PLACE1007282	8.6	4.51	8.76	10.51	12.35	10.29	22.66	27.14	27.14		**	+	
	PLACE1007286	6	1.42	3.35	6.08	8.09	5.91	3.36	4.27	4.27				
	PLACE1007296	5.96	3.96	4.56	9.09	9.08	8.48	6.51	8.92	8.92	**	+	*	+
	PLACE1007301	1.48	0.84	0.72	0.94	1.65	0.98	0.49	0.96	0.96				
	PLACE1007314	7.72	5.09	4.39	7.99	9.50	9.98	8.19	8.10	8.1	*	+		
35	PLACE1007317	1.71	0.70	0.71	2.11	1.11	1.58	1.38	1.29	1.29				
	PLACE1007329	1.19	1.05	0.73	3.19	2.34	1.79	1.73	2.65	2.65	*	+	*	+
	PLACE1007338	5.4	1.79	2.69	4.68	5.71	4.16	3.17	5.55	5.55				
	PLACE1007342	2.46	2.38	1.37	2.04	2.30	2.39	2.65	5.91	5.91				
	PLACE1007345	2.86	1.45	1.69	3.47	3.21	3.18	2.59	3.21	3.21	*	+		
	PLACE1007346	5.8	4.00	4.67	8.73	7.57	8.39	4.92	8.73	8.73	**	+		
40	PLACE1007359	3.11	1.64	2.21	3.58	2.56	2.94	3.24	3.82	3.82		*	+	
	PLACE1007367	9.92	5.57	5.83	12.43	19.19	16.79	8.33	10.26	10.26	*	+		
	PLACE1007375	1.77	1.76	1.63	2.23	2.83	2.75	1.31	0.63	0.63	*	+	*	-
	PLACE1007377	4.63	2.52	2.53	3.52	3.56	1.75	2.11	3.18	3.18				
	PLACE1007386	1.87	0.97	0.83	6.47	6.90	6.45	4.13	3.04	3.04	**	+	*	+
45	PLACE1007392	2.72	3.07	3.82	2.83	2.94	3.03	2.89	3.43	3.43				
	PLACE1007402	2.84	2.88	1.67	3.44	3.03	2.39	3.94	2.99	2.99				
	PLACE1007409	0.93	0.91	1.34	1.36	1.18	1.53	1.35	1.51	1.51				
	PLACE1007416	1.46	1.48	1.61	3.34	3.06	3.14	3.57	4.84	4.84	**	+	**	+
	PLACE1007420	9.86	15.04	12.94	15.1	14.93	22.60	15.85	14.30	14.3				
	PLACE1007431	0.76	1.71	1.22	1.19	2.25	1.99	1.64	1.51	1.51				
50	PLACE1007450	4.02	1.67	1.64	5.21	4.82	5.24	2.44	2.49	2.49	*	+		
	PLACE1007452	2.24	1.00	1.94	2.6	2.82	2.49	1.40	3.18	3.18				
	PLACE1007454	10.17	5.34	5.33	10.9	13.21	14.02	9.07	11.14	11.14	*	+		
	PLACE1007460	3.51	2.45	2.56	3.47	3.50	3.36	2.34	2.84	2.84				
	PLACE1007478	1.85	1.34	0.98	2.14	2.65	2.21	0.62	1.89	1.89	*	+		
55	PLACE1007484	1.62	2.03	1.82	4.03	4.26	4.32	4.30	4.48	4.48	**	+	**	+
	PLACE1007488	2.83	1.13	1.39	2.08	1.66	1.77	1.01	1.64	1.64				
	PLACE1007507	4.17	3.85	4.18	3.46	1.91	4.23	1.30	2.17	2.17		**	-	

Table 318

	PLACE1006711	7.17	2.48	3.66	6.98	7.47	5.78	4.03	4.95	4.95			
5	PLACE1006714	3.92	2.24	1.78	5.56	4.95	3.81	3.00	4.91	4.91			
	PLACE1006716	2.25	1.27	1.41	2.91	2.85	2.05	3.03	4.59	4.59		°	+
	PLACE1006731	2.78	1.41	1.10	2.51	2.88	3.14	3.12	3.70	3.7		°	+
	PLACE1006754	2.7	1.40	1.42	2.85	1.89	2.31	2.05	2.80	2.8			
	PLACE1006760	3.7	1.96	3.99	17.24	15.19	18.35	5.74	7.75	7.75	°°	+	°
10	PLACE1006779	0.53	0.60	0.34	1.36	0.57	1.21	0.75	1.01	1.01		°	+
	PLACE1006782	3.05	2.67	1.94	3.22	2.17	3.97	2.17	3.27	3.27			
	PLACE1006783	2.73	1.09	1.46	2.19	2.99	2.41	1.48	1.96	1.96			
	PLACE1006786	2.68	1.84	0.83	3.12	2.79	4.30	2.72	2.69	2.69			
	PLACE1006792	5.78	3.42	3.75	8.62	10.09	8.98	4.28	5.86	5.86	°°	+	
15	PLACE1006795	0.68	0.34	0.21	1.2	1.49	1.27	1.37	1.67	1.67	°°	+	°°
	PLACE1006810	0.58	0.50	0.45	1.01	1.36	1.09	0.49	1.98	1.98	°°	+	
	PLACE1006815	1.33	0.93	2.03	1.99	1.23	2.62	4.47	8.37	8.37		°	+
	PLACE1006819	3.99	2.53	2.85	4.94	4.18	4.26	2.87	3.81	3.81			
	PLACE1006815	2.42	2.62	2.14	3.2	3.02	2.39	2.60	2.42	2.42			
20	PLACE1006819	0.94	0.46	0.62	1.41	2.34	1.11	0.55	1.74	1.74			
	PLACE1006820	4.68	2.07	1.78	6.12	5.69	5.61	3.23	3.27	3.27	°	+	
	PLACE1006826	5.96	2.02	3.35	4.28	4.36	3.41	2.91	3.64	3.64			
	PLACE1006829	5.22	3.72	3.02	4.2	5.82	4.43	2.98	5.22	5.22			
	PLACE1006833	1.92	0.96	0.85	1.93	2.19	2.15	1.79	1.77	1.77			
	PLACE1006850	0.52	0.28	0.19	0.7	1.33	1.10	0.18	0.88	0.88	°	+	
25	PLACE1006857	3.61	1.51	1.29	3.02	3.99	3.62	1.66	1.92	1.92			
	PLACE1006878	3.81	2.86	3.20	2.81	3.41	2.95	2.46	3.28	3.28			
	PLACE1006878	2.74	2.03	2.05	2.44	3.93	2.25	1.87	2.15	2.15			
	PLACE1006883	6.43	2.64	2.47	5.83	6.59	4.26	4.67	3.84	3.84			
	PLACE1006890	2.65	0.75	0.60	1.14	1.52	1.02	0.75	1.07	1.07			
30	PLACE1006901	2.51	0.47	1.17	2.93	3.57	2.34	0.90	1.69	1.69			
	PLACE1006924	2.19	1.14	0.97	3.15	2.91	3.59	2.13	2.06	2.06	°	+	
	PLACE1006917	6.14	2.79	3.06	4.32	4.29	4.20	3.17	2.44	2.44			
	PLACE1006932	5	1.78	2.39	3.19	3.17	4.46	2.94	4.82	4.82			
	PLACE1006935	2.14	0.74	0.92	1.51	0.93	2.00	1.13	1.70	1.7			
35	PLACE1006936	4.8	2.30	2.67	3.82	4.93	3.67	2.67	3.02	3.02			
	PLACE1006938	3.3	0.68	0.97	1.15	2.53	1.83	2.18	2.76	2.76			
	PLACE1006939	5.12	2.95	4.08	5.45	7.11	5.94	4.25	6.06	6.06			
	PLACE1006951	6.24	3.14	3.71	8.87	11.45	12.47	5.75	6.96	6.96	°	+	
	PLACE1006952	3.09	1.63	2.08	6.06	7.00	5.67	3.12	4.82	4.82	°°	+	°
	PLACE1006956	3.67	1.18	1.70	1.85	1.83	1.79	1.92	2.51	2.51			
40	PLACE1006979	2	0.97	1.09	2.59	1.79	2.03	1.44	1.20	1.2			
	PLACE1006989	6.78	4.06	4.71	5.85	5.19	8.95	4.33	4.95	4.95			
	PLACE1007001	4.54	2.23	1.52	6.32	8.61	5.77	3.73	6.03	6.03	°	+	
	PLACE1007014	7.18	3.58	3.26	4.66	5.59	4.03	3.90	5.33	5.33			
	PLACE1007021	1.97	0.96	1.13	2.46	2.25	1.64	1.52	0.94	0.94			
45	PLACE1007026	2.03	0.23	0.75	2.47	2.67	2.53	4.15	4.32	4.32	°	+	°°
	PLACE1007028	3.59	1.48	2.53	3.68	2.34	2.63	3.78	4.37	4.37			
	PLACE1007038	9.6	3.28	7.64	12.57	9.19	16.41	73.23	81.92	81.92		°°	+
	PLACE1007040	3.28	1.64	2.20	3.38	2.82	2.64	2.43	2.13	2.13			
	PLACE1007045	2.23	0.95	1.52	6.73	6.04	4.31	5.12	5.62	5.62	°°	+	°°
50	PLACE1007045	283.34	168.88	128.09	131.34	214.44	117.39	119.27	112.98	113			
	PLACE1007053	5.82	1.54	2.58	3.59	4.38	2.76	2.77	4.36	4.36			
	PLACE1007059	5.93	3.20	2.64	4.13	5.60	3.12	3.77	3.46	3.46			
	PLACE1007070	1.79	1.14	1.74	2.68	3.48	2.65	2.23	3.69	3.69	°	+	°
	PLACE1007076	49.7	17.82	25.75	20.08	16.26	21.00	15.39	17.24	17.24			
55	PLACE1007077	2.93	1.23	2.63	2.96	2.01	1.85	3.14	3.21	3.21			
	PLACE1007081	1.29	0.25	0.75	1.37	1.15	1.43	0.54	1.23	1.23			
	PLACE1007082	8.76	4.12	5.94	5.68	4.75	5.79	2.91	3.11	3.11			

Table 317

5	PLACE1006368	8.01	4.14	3.31	4.49	6.87	3.91	2.14	3.48	3.48				
	PLACE1006371	3.39	1.39	1.67	3.81	5.96	2.01	3.24	1.56	1.56				
	PLACE1006373	3.53	2.18	2.19	3.47	4.18	3.83	2.92	2.88	2.88				
	PLACE1006382	0.97	0.61	1.44	1.43	2.65	2.53	1.94	2.62	2.62		*	+	
	PLACE1006385	4.48	1.74	2.64	3.36	3.37	3.94	3.78	4.13	4.13				
10	PLACE1006391	2.37	0.62	1.55	2.01	1.29	1.94	1.72	2.56	2.56				
	PLACE1006412	4.8	2.68	3.85	7.96	8.66	10.16	7.60	4.54	4.54	**	+		
	PLACE1006414	1.25	0.89	0.94	1.45	2.86	1.96	0.92	1.08	1.08				
	PLACE1006419	17.56	9.39	8.08	6.95	7.32	5.48	8.27	8.11	8.11				
	PLACE1006438	8.55	3.61	3.22	5.14	6.25	6.01	5.43	4.95	4.95				
	PLACE1006443	13.27	8.13	8.94	9.9	11.09	10.15	9.09	10.58	10.58				
15	PLACE1006445	4.37	2.38	3.95	6.95	9.30	6.55	3.68	5.38	5.38	*	+		
	PLACE1006447	3.95	1.73	1.16	4.37	4.04	4.18	2.52	2.55	2.55				
	PLACE1006466	2.16	1.21	1.47	2	2.00	2.12	1.67	2.19	2.19				
	PLACE1006469	5.27	2.73	2.42	5.93	3.56	4.11	2.77	4.56	4.56				
	PLACE1006470	5.41	1.20	2.14	5.2	5.53	6.27	4.08	3.01	3.01				
20	PLACE1006472	11.56	7.21	5.05	18.35	19.64	11.78	13.72	15.01	15.01	*	+	*	+
	PLACE1006476	5.69	2.73	2.21	5.81	8.49	6.21	4.48	5.62	5.62				
	PLACE1006482	2.17	1.70	2.74	3.32	3.51	3.07	2.54	2.44	2.44	*	+		
	PLACE1006488	12.25	5.32	6.03	9.43	11.28	10.04	10.74	9.34	9.34				
	PLACE1006492	6.49	3.62	3.60	9.32	9.53	11.55	11.09	11.09	11.09	**	+	**	+
	PLACE1006506	4.02	1.67	1.46	3.66	1.98	4.89	2.21	2.62	2.62				
25	PLACE1006515	1.42	1.65	2.04	2.45	1.89	3.92	0.81	1.40	1.4				
	PLACE1006516	2.44	0.98	1.54	4.26	3.82	5.07	3.64	3.02	3.02	**	+	*	+
	PLACE1006520	3.63	0.73	1.91	3.9	6.61	4.44	1.81	3.39	3.39				
	PLACE1006521	6.56	3.47	2.11	9.33	11.45	8.09	6.98	6.31	6.31	*	+		
	PLACE1006529	8.21	3.84	3.76	6.99	8.95	8.26	5.00	11.36	11.36				
30	PLACE1006531	4.94	2.43	2.89	5.42	4.81	4.48	4.13	3.68	3.68				
	PLACE1006534	5.02	1.96	2.25	4.42	4.01	5.10	4.71	2.91	2.91				
	PLACE1006540	7.85	3.19	3.56	8.91	8.99	10.06	5.53	6.70	6.7	*	+		
	PLACE1006549	6.58	4.45	4.11	5.8	5.03	4.33	3.92	6.01	6.01				
	PLACE1006550	5.23	2.28	2.45	4.69	4.00	3.88	3.29	3.49	3.49				
35	PLACE1006552	6.12	1.72	2.67	5.75	4.74	3.07	2.71	2.86	2.86				
	PLACE1006557	5.34	2.94	3.14	4.05	3.81	4.16	3.41	4.94	4.94				
	PLACE1006563	9.2	2.53	5.98	6.32	8.19	6.80	4.10	7.57	7.57				
	PLACE1006579	2.63	1.19	1.62	2.98	3.80	3.82	2.66	2.84	2.84	*	+		
	PLACE1006594	2.07	1.44	0.90	5.07	5.06	4.32	1.36	3.33	3.33	**	+		
40	PLACE1006598	1.81	0.42	0.76	1.91	2.22	2.18	1.31	2.09	2.09				
	PLACE1006607	3.34	1.19	1.08	3.9	3.89	4.86	2.07	2.61	2.61	*	+		
	PLACE1006610	8.31	5.63	5.00	11.87	9.53	10.14	8.32	7.46	7.46	*	+		
	PLACE1006615	14.76	9.42	9.72	14.75	13.78	11.87	9.86	12.58	12.58				
	PLACE1006617	3.05	1.29	1.68	3.75	3.86	3.39	2.76	2.76	2.76	*	+		
	PLACE1006618	6.92	2.44	3.52	4.27	4.86	5.91	4.69	6.70	6.7				
45	PLACE1006626	5.11	2.06	2.30	4.94	4.91	4.81	2.78	5.05	5.05				
	PLACE1006629	0.66	0.42	0.61	1.08	1.24	1.75	1.19	2.37	2.37	*	+	*	+
	PLACE1006637	4.27	1.61	1.80	4.69	6.26	5.06	2.25	2.76	2.76	*	+		
	PLACE1006640	0.61	0.64	0.44	0.58	0.66	0.88	0.53	0.93	0.93				
	PLACE1006644	4.05	2.79	2.37	3.98	3.74	4.19	3.10	4.40	4.4				
50	PLACE1006657	2	0.91	0.90	4.27	3.82	3.51	2.62	2.98	2.98	**	+	*	+
	PLACE1006673	4.86	2.75	3.39	6.61	5.20	7.30	3.73	3.71	3.71	*	+		
	PLACE1006678	2.03	0.79	2.52	1.93	2.13	1.51	1.82	3.83	3.83				
	PLACE1006682	12.66	6.71	7.70	9.17	11.56	9.08	12.78	10.93	10.93				
	PLACE1006684	0.85	0.40	0.58	0.51	0.62	0.80	0.74	0.98	0.98				
	PLACE1006698	2.49	1.60	2.01	2.82	3.08	3.05	2.36	3.97	3.97	*	+		
55	PLACE1006704	2.61	1.71	2.74	4.42	5.65	7.34	3.32	3.83	3.83	*	+	*	+
	PLACE1006708	5.71	3.09	2.56	5.97	10.34	7.12	1.92	5.99	5.99				

Table 316

5	PLACE1006002	8.53	4.41	5.77	19.13	16.78	16.67	8.23	9.69	9.69	**	+		
	PLACE1006003	6.88	5.62	5.05	3.42	5.00	5.45	4.05	7.43	7.43				
	PLACE1006011	4.72	2.78	3.04	3.63	3.41	3.26	2.90	3.61	3.61				
	PLACE1006017	4.17	1.57	1.37	3.12	3.78	3.87	3.13	4.29	4.29				
	PLACE1006037	8.36	3.71	4.44	4.09	4.76	4.29	2.99	4.73	4.73				
10	PLACE1006040	13.34	8.65	10.10	9.09	7.82	11.18	9.13	10.46	10.46				
	PLACE1006063	4.18	2.39	2.46	2.52	3.00	2.07	2.59	2.91	2.91				
	PLACE1006071	3.1	2.05	2.07	1.68	2.75	3.43	1.83	2.76	2.76				
	PLACE1006073	3.97	2.14	1.81	6.25	6.16	5.43	3.65	5.10	5.1	*	+		
	PLACE1006074	4.44	2.36	2.42	6.36	6.76	5.83	2.98	4.13	4.13	*	+		
	PLACE1006076	1.24	0.92	1.14	3.37	4.38	2.74	2.16	3.59	3.59	**	+	*	+
15	PLACE1006079	4.64	2.47	2.65	3.89	4.84	4.04	4.58	5.85	5.85				
	PLACE1006093	1.06	0.90	1.72	1.34	1.63	0.86	2.10	2.38	2.38		*	+	
	PLACE1006116	2.79	1.95	1.97	2.66	2.53	2.69	3.38	3.33	3.33		*	+	
	PLACE1006119	2.59	2.94	2.87	5.28	4.68	6.57	3.23	3.84	3.84	**	+	*	+
	PLACE1006129	2.82	1.25	0.50	2.84	2.73	3.10	3.07	1.53	1.53				
20	PLACE1006139	7.84	6.54	4.25	6.48	5.34	5.86	6.94	4.78	4.78				
	PLACE1006143	2.36	1.84	1.60	4.6	3.86	4.22	1.68	3.18	3.18	**	+		
	PLACE1006157	2.84	1.26	1.64	2.25	2.35	1.82	1.52	2.36	2.36				
	PLACE1006159	1.74	1.38	1.27	2.48	3.25	2.76	3.72	4.61	4.61	**	+	**	+
	PLACE1006164	0.77	0.31	0.34	1.19	1.94	1.20	1.01	0.99	0.99	*	+	*	+
25	PLACE1006167	6.97	5.82	7.53	6.63	9.38	9.10	8.80	7.88	7.88				
	PLACE1006170	3.23	2.05	2.23	3.8	5.15	4.56	3.39	4.89	4.89	*	+	*	+
	PLACE1006181	4.1	2.72	3.53	6.41	6.16	6.21	5.86	6.48	6.48	**	+	**	+
	PLACE1006187	0.5	0.33	0.10	0.86	0.82	1.09	0.66	0.49	0.49	*	+		
	PLACE1006195	3.24	1.23	1.17	2.67	2.87	2.14	2.62	1.30	1.3				
	PLACE1006196	8.03	2.93	3.80	5.31	7.47	6.96	4.75	3.79	3.79				
30	PLACE1006197	7.57	3.83	6.49	6.35	7.27	5.99	3.44	4.86	4.86				
	PLACE1006198	2.55	1.19	1.79	2.81	2.56	2.19	0.91	2.46	2.46				
	PLACE1006205	0.84	0.89	1.05	0.57	0.49	1.57	0.74	1.36	1.36				
	PLACE1006208	2.19	1.80	3.16	5	4.18	5.05	7.99	4.42	4.42	**	+		
	PLACE1006211	24.46	16.10	17.64	12.62	6.69	13.24	6.25	5.01	5.01		**	-	
35	PLACE1006219	3.37	2.25	3.36	4.14	6.29	3.89	6.74	5.53	5.53		**	+	
	PLACE1006223	1.64	1.06	2.11	4.34	4.03	4.26	2.35	1.45	1.45	**	+		
	PLACE1006225	1.79	1.20	1.26	2	1.95	1.83	1.23	1.27	1.27				
	PLACE1006236	1.44	1.01	1.87	3.01	4.09	1.96	1.59	2.02	2.02				
	PLACE1006239	1.72	1.00	1.18	2.46	2.48	2.60	1.22	3.24	3.24	**	+		
40	PLACE1006245	3.4	2.04	2.24	3.29	3.77	3.95	2.39	2.28	2.28				
	PLACE1006246	2.78	1.91	2.09	2.77	4.35	3.44	2.43	1.97	1.97				
	PLACE1006248	1.93	1.11	1.30	3.09	2.89	3.27	2.22	1.63	1.63	**	+		
	PLACE1006262	3.84	0.83	1.42	2.66	2.88	1.75	2.15	1.21	1.21				
	PLACE1006269	3.04	1.04	0.97	1.76	1.81	1.87	1.28	1.88	1.88				
45	PLACE1006275	7.22	3.03	3.50	4.21	3.01	5.50	3.90	5.68	5.68				
	PLACE1006277	2.96	1.17	2.13	3.73	2.55	2.62	1.86	2.11	2.11				
	PLACE1006288	11.06	5.08	6.39	7.42	7.01	7.05	7.31	9.33	9.33				
	PLACE1006290	2.57	0.88	2.08	1.76	1.36	2.22	3.49	3.56	3.56		*	+	
	PLACE1006298	4.88	1.93	2.02	5.27	5.25	6.12	2.74	3.06	3.06				
	PLACE1006311	0.92	0.15	0.41	0.82	0.31	0.81	2.11	2.64	2.64		**	+	
50	PLACE1006318	4.74	1.72	2.13	3.2	4.07	2.67	3.29	3.63	3.63				
	PLACE1006325	9.29	2.77	3.97	3.52	11.00	5.12	4.02	5.45	5.45				
	PLACE1006331	4.1	2.50	3.35	6.3	7.84	7.16	3.59	5.15	5.15	**	+		
	PLACE1006335	4.07	1.71	1.37	3.4	2.75	3.34	2.92	1.94	1.94				
	PLACE1006357	1.19	0.25	0.48	0.63	0.90	0.73	0.68	1.54	1.54				
55	PLACE1006360	5.46	2.85	2.81	3.84	5.35	5.87	2.18	2.76	2.76				
	PLACE1006364	2.49	1.09	1.52	2.96	1.99	3.19	1.91	2.29	2.29				
	PLACE1006365	0.49	0.34	1.09	1.19	1.22	2.88	0.94	1.48	1.48				

Table 315

5	PLACE1005682	2.11	2.05	2.13	4.34	3.23	4.41	1.89	2.15	2.15	**	+		
	PLACE1005698	4.64	2.14	-3.28	3.89	3.92	4.16	1.91	2.53	2.53				
	PLACE1005708	25.78	13.70	10.51	13.88	16.18	11.27	14.00	14.43	14.43				
	PLACE1005725	3.83	1.42	2.33	2.34	3.92	2.04	4.70	4.61	4.61		*	+	
	PLACE1005727	8.48	2.60	3.97	5.4	4.41	4.96	2.49	2.57	2.57				
10	PLACE1005730	3.57	0.90	1.62	1.95	2.02	2.00	2.05	2.95	2.95				
	PLACE1005736	4.39	2.36	2.88	8.34	10.28	9.63	5.13	7.81	7.81	**	+	*	+
	PLACE1005739	2.31	1.03	1.11	1.47	1.17	1.64	2.22	2.15	2.15				
	PLACE1005745	9.25	5.63	5.40	10.32	14.44	8.66	7.38	8.69	8.69				
	PLACE1005752	4.63	2.11	0.91	2.57	2.97	2.88	2.25	2.86	2.86				
15	PLACE1005755	0.83	0.18	0.42	0.66	1.88	0.66	0.70	0.93	0.93				
	PLACE1005756	14.63	7.31	9.39	22.2	25.42	27.72	29.92	35.68	35.68	**	+	**	+
	PLACE1005760	7.89	3.72	4.80	10.59	12.05	10.96	9.45	9.92	9.92	*	+	*	+
	PLACE1005763	3.86	1.70	3.26	6.59	6.36	6.88	4.43	4.28	4.28	**	+		
	PLACE1005768	6.14	3.01	5.24	7.97	7.90	8.87	6.22	5.90	5.9	*	+		
	PLACE1005771	7.62	3.12	5.03	7.4	7.32	9.76	6.04	6.48	6.48				
20	PLACE1005783	3.63	1.45	2.35	2.79	4.79	2.04	2.34	3.07	3.07				
	PLACE1005799	6.45	3.16	3.38	5.32	4.64	3.49	5.15	5.23	5.23				
	PLACE1005802	5.01	1.66	1.63	4.46	8.45	4.41	2.49	4.79	4.79				
	PLACE1005803	11.48	4.59	6.77	9.23	10.65	9.39	6.53	8.91	8.91				
	PLACE1005804	1.62	0.72	0.84	1.97	2.36	1.93	2.21	2.56	2.56	*	+	*	+
25	PLACE1005813	10.74	3.23	5.61	11.66	8.19	9.55	6.52	6.57	6.57				
	PLACE1005815	5.12	2.48	3.85	7.34	9.35	11.87	4.89	5.17	5.17	*	+		
	PLACE1005828	5.16	3.37	3.80	8.35	8.98	9.59	4.86	6.29	6.29	**	+		
	PLACE1005833	3.06	1.35	1.59	18.69	21.23	11.91	28.00	30.88	30.88	**	+	**	+
	PLACE1005834	1.93	0.65	0.55	4	6.43	2.66	1.50	2.50	2.5	*	+		
30	PLACE1005835	5.07	4.66	2.88	5.05	7.51	3.87	4.83	4.52	4.52				
	PLACE1005836	3.75	1.63	2.11	2.62	6.42	3.23	2.73	2.06	2.06				
	PLACE1005845	4.98	1.86	2.24	4.26	4.56	2.61	2.60	3.15	3.15				
	PLACE1005850	4.23	2.74	2.58	5.55	4.59	5.10	2.95	3.19	3.19	*	+		
	PLACE1005851	1.83	0.96	1.69	2.54	2.84	4.11	1.02	0.85	0.85	*	+		
	PLACE1005856	4.08	1.29	7.53	4.1	2.89	3.39	1.78	2.05	2.05				
35	PLACE1005875	3.56	1.05	0.65	5.19	5.82	3.59	3.48	3.10	3.1				
	PLACE1005876	4.08	3.91	2.72	2.79	2.82	2.10	2.04	2.27	2.27		*	-	
	PLACE1005878	5.27	2.13	2.19	4.92	3.53	2.84	3.83	3.82	3.82				
	PLACE1005880	3.44	0.96	1.32	2.14	2.64	2.46	2.97	4.34	4.34				
	PLACE1005884	1.76	0.52	0.55	1.39	1.77	1.41	2.43	2.29	2.29		*	+	
40	PLACE1005890	2.04	0.70	0.65	1.41	1.86	1.52	1.88	2.21	2.21				
	PLACE1005898	2.99	2.09	1.71	4.94	3.42	2.88	2.46	3.27	3.27				
	PLACE1005913	5.71	2.57	3.76	7.83	8.39	8.51	3.79	4.62	4.62	*	+		
	PLACE1005921	10.98	4.34	4.34	9.34	8.32	8.81	6.16	6.43	6.43				
	PLACE1005923	57.96	26.97	25.39	4.09	4.25	2.49	3.95	3.48	3.48	*	-	*	-
45	PLACE1005925	2.51	0.91	2.14	3.11	3.71	2.82	1.93	2.80	2.8				
	PLACE1005927	6.09	2.70	1.89	3.69	4.68	4.01	3.18	5.73	5.73				
	PLACE1005932	1.82	0.71	0.41	1.33	1.76	1.08	1.15	1.54	1.54				
	PLACE1005934	3.84	2.41	2.72	6.26	7.11	6.43	3.93	5.30	5.3	**	+	*	+
	PLACE1005936	2.29	0.78	1.05	1.64	1.78	1.51	1.47	1.36	1.36				
	PLACE1005939	6.69	4.35	3.92	5.44	5.46	3.73	16.40	25.30	25.3		**	+	
50	PLACE1005951	5.63	2.39	2.32	4.86	4.02	3.74	3.42	3.43	3.43				
	PLACE1005953	2.92	1.24	0.96	2.51	1.78	2.30	1.15	2.55	2.55				
	PLACE1005955	3.7	1.62	1.62	2.35	3.61	2.40	3.03	3.07	3.07				
	PLACE1005966	3.38	0.65	1.08	1.86	2.32	2.12	2.02	2.80	2.8				
	PLACE1005968	10.55	5.64	5.72	7.48	5.88	6.36	5.68	7.85	7.85				
55	PLACE1005975	10.44	5.19	8.44	12.95	14.63	15.64	8.66	16.92	16.92	*	+		
	PLACE1005990	3.19	1.43	1.58	2.28	2.05	1.79	1.80	3.26	3.26				
	PLACE1005997	64.81	36.05	40.42	54.4	53.64	53.12	27.58	33.55	33.55				

Table 314

5	PLACE1005374	5	2.10	2.77	8.04	11.61	11.01	4.31	6.01	6.01	**	+		
	PLACE1005383	8.86	3.18	3.37	5.63	6.03	4.19	5.25	6.23	6.23				
	PLACE1005388	2.57	0.54	0.31	2.75	1.56	0.89	2.61	1.22	1.22				
	PLACE1005409	5.48	3.06	2.63	7.59	8.06	6.25	3.31	4.02	4.02	*	+		
	PLACE1005410	6.76	2.97	3.65	5.66	8.24	5.17	9.00	11.77	11.77		*	+	
	PLACE1005426	4.46	1.72	1.45	2.27	1.48	1.00	3.43	3.54	3.54				
10	PLACE1005431	4.56	2.63	2.58	4.42	5.14	6.40	5.57	6.50	6.5		*	+	
	PLACE1005453	3.55	1.77	2.09	4.33	4.49	5.14	1.74	3.20	3.2	*	+		
	PLACE1005467	5.64	2.78	2.70	6.57	5.73	4.48	5.05	4.51	4.51				
	PLACE1005471	3.36	0.50	1.20	3.42	3.09	2.65	2.30	3.64	3.64				
	PLACE1005476	5.15	1.54	1.43	2.43	2.59	1.89	1.59	3.01	3.01				
15	PLACE1005477	2.24	1.35	1.27	5.66	7.05	5.00	4.23	7.05	7.05	**	+	*	+
	PLACE1005480	1.93	1.39	1.29	1.24	1.52	1.24	1.31	1.75	1.75				
	PLACE1005481	2.22	1.41	1.51	2.73	2.46	3.04	1.87	2.00	2	*	+		
	PLACE1005494	1.24	0.38	0.90	0.8	0.90	0.66	0.80	1.98	1.98				
	PLACE1005495	4.56	1.60	1.71	3.4	2.67	2.72	2.06	1.93	1.93				
20	PLACE1005497	8.06	4.83	3.69	4.42	2.88	4.07	9.50	10.40	10.4		*	+	
	PLACE1005499	4.76	1.36	1.66	2.69	4.07	3.13	5.56	5.51	5.51				
	PLACE1005502	2.69	0.87	1.10	2.75	3.41	2.24	1.89	4.02	4.02				
	PLACE1005513	1.27	0.71	0.80	3.5	2.88	3.38	1.95	3.18	3.18	**	+	*	+
	PLACE1005515	2.84	0.81	0.90	1.12	0.96	1.43	2.38	3.90	3.9				
25	PLACE1005519	7.14	2.92	5.14	2.37	3.46	3.11	2.55	3.35	3.35				
	PLACE1005526	2.06	1.07	1.41	1.41	2.39	1.85	1.31	2.23	2.23				
	PLACE1005528	6.82	2.99	3.77	7.7	10.09	11.05	4.64	5.96	5.96	*	+		
	PLACE1005530	4.98	2.54	2.80	2.85	5.04	3.55	3.48	2.83	2.83				
	PLACE1005536	4.27	3.13	1.98	6.1	4.77	1.67	4.10	3.87	3.87				
	PLACE1005539	3	1.66	1.31	3.17	3.20	2.66	1.69	3.05	3.05				
30	PLACE1005543	2.3	1.25	1.18	4	3.96	4.38	3.55	3.32	3.32	**	+	**	+
	PLACE1005544	6.06	3.23	2.89	3.81	4.11	4.35	4.12	5.12	5.12				
	PLACE1005550	8.49	4.71	5.86	4.53	4.75	4.40	2.14	3.57	3.57				
	PLACE1005554	1.55	0.76	0.94	1.77	1.45	1.38	2.99	1.56	1.56				
	PLACE1005557	3.3	1.97	2.34	3.4	5.03	3.76	3.56	3.17	3.17				
35	PLACE1005563	1.99	2.09	0.76	1.69	2.10	1.89	2.11	1.69	1.69				
	PLACE1005569	4.54	2.73	2.52	4.62	4.22	2.24	2.63	3.22	3.22				
	PLACE1005574	1.43	0.92	0.87	2.29	2.41	2.10	0.45	0.99	0.99	**	+		
	PLACE1005584	1.32	0.88	0.93	1.31	1.40	1.67	1.68	4.67	4.67				
	PLACE1005590	2.53	3.81	2.63	3.18	2.75	3.39	4.08	5.93	5.93		*	+	
	PLACE1005595	2.91	2.55	3.00	2.96	2.39	3.53	3.75	3.64	3.64		**	+	
40	PLACE1005601	2.77	1.99	2.02	2.52	2.79	3.50	2.97	3.86	3.86		*	+	
	PLACE1005603	0.9	0.55	0.69	0.87	1.06	0.76	1.27	1.79	1.79		*	+	
	PLACE1005604	4.18	2.56	1.82	4.89	4.83	6.27	2.39	1.93	1.93	*	+		
	PLACE1005611	2.64	2.26	1.19	5.02	2.53	3.51	2.64	2.53	2.53				
	PLACE1005622	2.15	1.96	1.00	2.49	2.91	2.25	1.48	2.00	2				
45	PLACE1005623	4.29	1.35	2.10	3.3	3.81	3.92	2.17	2.70	2.7				
	PLACE1005630	6.26	3.63	2.27	4.66	6.06	5.41	4.45	5.87	5.87				
	PLACE1005639	1.47	1.40	2.08	1.45	2.40	2.84	0.78	1.78	1.78				
	PLACE1005646	5.91	4.51	5.24	4.63	5.74	5.46	4.51	5.47	5.47				
	PLACE1005647	0.51	0.39	0.52	1.16	1.74	1.90	2.41	4.04	4.04	**	+	**	+
	PLACE1005648	5.72	4.93	6.25	15.18	16.23	16.45	5.58	7.21	7.21	**	+		
50	PLACE1005653	3.3	1.90	0.82	3.94	5.11	4.03	2.85	2.07	2.07	*	+		
	PLACE1005656	2.07	1.04	0.59	1.23	2.09	0.91	1.88	0.45	0.45				
	PLACE1005659	4.14	1.56	2.46	2.97	4.17	2.22	1.44	2.31	2.31				
	PLACE1005660	5.27	3.90	2.60	4.31	5.01	2.96	3.33	4.29	4.29				
	PLACE1005664	4.13	4.07	4.07	5.57	5.47	4.07	5.14	6.25	6.25			**	+
55	PLACE1005666	0.97	1.45	1.51	3.22	3.91	4.93	3.26	2.77	2.77	**	+	**	+
	PLACE1005669	4.53	2.92	2.87	6.24	4.95	7.16	3.36	4.69	4.69	*	+		

Table 313

5	PLACE1005055	1.93	1.90	2.25	2.55	3.80	3.83	1.39	2.30	2.3	*	+		
	PLACE1005066	3.73	3.53	2.95	3.62	2.74	3.71	4.65	6.92	6.92		*	+	
	PLACE1005077	1.88	0.74	0.51	1.94	2.30	1.62	1.19	1.27	1.27				
	PLACE1005085	5.35	2.26	1.94	7.82	9.01	6.89	4.04	4.10	4.1	*	+		
	PLACE1005086	8.18	4.09	4.61	8.82	11.72	8.88	4.94	5.91	5.91				
10	PLACE1005088	48.83	27.68	29.69	27.61	39.82	34.65	26.01	25.68	25.68				
	PLACE1005089	2.42	1.38	1.99	2.77	2.07	2.49	2.33	3.56	3.56				
	PLACE1005101	6.75	6.64	8.03	8.45	9.96	12.39	8.67	10.11	10.11		*	+	
	PLACE1005102	5.88	7.51	8.49	11.05	10.78	12.60	9.73	9.59	9.59	*	+	*	+
	PLACE1005108	5.63	4.27	3.64	12.01	12.87	10.10	5.64	5.46	5.46	**	+		
15	PLACE1005110	6.84	3.16	2.29	5.61	4.42	2.27	2.47	3.96	3.96				
	PLACE1005111	2.32	1.43	0.52	2.8	3.48	1.64	1.69	1.48	1.48				
	PLACE1005123	20.53	8.57	10.06	12.54	14.07	10.45	7.24	8.30	8.3				
	PLACE1005124	3.92	2.40	2.02	3.08	6.72	4.08	3.28	3.46	3.46				
	PLACE1005128	10.6	9.42	9.74	12.9	15.61	15.03	14.09	17.89	17.89	**	+	**	+
	PLACE1005130	4.63	4.42	3.58	6.21	6.12	6.60	2.90	3.62	3.62	**	+		
20	PLACE1005141	11.53	6.88	7.85	10.2	11.46	13.07	6.08	6.65	6.65				
	PLACE1005146	2.66	2.45	2.31	3.79	4.23	2.90	1.91	2.35	2.35	*	+		
	PLACE1005152	4.31	1.32	1.78	5.23	4.05	4.11	2.87	2.37	2.37				
	PLACE1005157	3.17	1.71	2.58	3.61	2.97	3.04	1.83	2.24	2.24				
	PLACE1005162	5.03	1.44	2.16	4.55	5.47	5.51	3.63	3.97	3.97				
25	PLACE1005170	1.73	0.31	0.62	1.61	1.26	1.41	1.34	1.72	1.72				
	PLACE1005176	1.61	0.38	0.68	1.16	1.34	1.12	1.06	1.60	1.6				
	PLACE1005181	0.5	0.24	0.53	1.19	0.87	2.59	0.77	1.26	1.26		*	+	
	PLACE1005184	4.44	1.78	2.90	7.9	7.10	9.09	4.75	4.64	4.64	**	+		
	PLACE1005186	6.95	2.41	3.82	3.37	3.80	2.87	3.22	3.68	3.68				
30	PLACE1005187	3.14	1.53	1.03	3.09	5.30	4.21	2.97	2.82	2.82				
	PLACE1005189	5.93	2.53	2.32	3.58	5.81	4.44	5.57	5.74	5.74				
	PLACE1005193	6.13	3.49	3.63	4.29	4.51	4.47	3.64	4.00	4				
	PLACE1005200	4.37	1.39	2.33	2.59	3.60	1.69	2.29	2.95	2.95				
	PLACE1005206	2.34	0.51	1.37	1.54	2.19	3.01	1.80	1.98	1.98				
35	PLACE1005216	1.38	0.71	1.11	2.26	2.41	2.76	2.43	3.73	3.73	**	+	**	+
	PLACE1005223	4.29	2.34	2.64	6.04	7.76	7.97	4.06	6.10	6.1	**	+		
	PLACE1005225	19.66	8.09	9.52	16.05	21.00	13.76	8.27	9.44	9.44				
	PLACE1005232	8.02	4.04	2.69	6.94	10.56	7.61	5.96	6.58	6.58				
	PLACE1005239	5.38	1.20	2.07	5.01	3.78	2.93	2.36	3.31	3.31				
	PLACE1005243	5.32	3.76	4.72	5.19	5.09	5.33	3.34	5.82	5.82				
40	PLACE1005250	3.75	1.12	1.85	3.16	3.89	3.16	2.16	2.84	2.84				
	PLACE1005261	2.07	0.70	1.90	2.25	2.05	1.77	2.13	1.93	1.93				
	PLACE1005266	1.9	0.95	1.09	2.57	2.39	2.64	2.14	1.90	1.9	*	+		
	PLACE1005271	5.66	2.63	3.94	8.71	9.11	8.37	4.71	5.02	5.02	**	+		
	PLACE1005277	3.05	0.82	0.70	2.46	4.32	1.50	1.02	2.07	2.07				
45	PLACE1005287	6.59	3.30	3.94	10.35	15.42	7.57	8.69	8.45	8.45		*	+	
	PLACE1005299	22.18	11.98	9.53	18.56	24.11	17.96	21.90	22.45	22.45				
	PLACE1005305	5.96	2.44	4.52	8.17	10.96	9.42	8.88	11.22	11.22	*	+	**	+
	PLACE1005307	3.74	1.42	2.86	4.85	5.32	3.53	2.69	4.11	4.11				
	PLACE1005308	3.94	1.81	2.45	3.16	2.71	2.64	2.67	2.60	2.6				
50	PLACE1005313	1.8	1.22	2.93	1.89	0.89	2.76	1.70	1.69	1.69				
	PLACE1005320	2.05	0.78	1.58	1.96	1.63	3.04	1.42	1.54	1.54				
	PLACE1005327	3.57	2.45	2.12	2.64	6.29	3.81	4.41	6.45	6.45		*	+	
	PLACE1005331	4	2.27	3.11	3.34	6.04	3.03	3.28	2.86	2.86				
	PLACE1005335	9.31	5.05	4.18	8.68	7.24	5.98	5.53	6.95	6.95				
	PLACE1005336	3.13	1.45	2.61	5.52	6.69	4.80	4.00	4.81	4.81	*	+	*	+
55	PLACE1005351	30.75	16.28	19.31	14.85	14.56	18.13	32.39	30.68	30.68				
	PLACE1005366	3.38	2.74	2.56	10.21	9.37	10.62	9.15	9.50	9.5	**	+	**	+
	PLACE1005373	4.26	1.58	2.70	3.39	2.69	4.82	2.63	3.29	3.29				

Table 312

5	PLACE1004743	2.83	1.69	1.62	2.65	2.30	3.35	1.64	2.59	2.59			
	PLACE1004751	3.88	2.76	2.71	4.3	4.81	6.07	2.16	4.03	4.03	*	+	
	PLACE1004757	6.62	2.79	3.38	5.64	5.36	5.13	4.59	3.33	3.33			
	PLACE1004761	1.53	0.69	0.99	1.89	2.90	1.43	1.17	2.01	2.01			
	PLACE1004773	6.07	1.81	3.15	5.28	4.05	5.04	3.00	3.37	3.37			
	PLACE1004775	0.59	0.48	0.41	0.54	0.33	0.45	0.35	1.11	1.11			
10	PLACE1004777	2.87	1.56	1.63	3.6	3.28	3.27	3.12	2.18	2.18	*	+	
	PLACE1004793	1.91	0.67	0.75	1.6	1.01	2.08	1.33	1.74	1.74			
	PLACE1004796	11.15	4.76	6.53	15.2	11.67	18.12	12.53	11.15	11.15	*	+	
	PLACE1004804	2.49	2.83	3.47	3.45	3.93	5.58	2.84	4.15	4.15			
	PLACE1004813	1.83	1.78	1.19	2.06	4.34	2.04	2.93	2.61	2.61		**	+
15	PLACE1004814	15.6	8.20	7.30	20.97	26.56	22.14	11.65	11.36	11.36	*	+	
	PLACE1004815	2.09	1.04	1.32	4.73	4.30	3.56	2.27	2.36	2.36	**	+	
	PLACE1004816	3.22	1.11	2.11	2.58	2.27	3.19	1.56	4.07	4.07			
	PLACE1004824	10.16	4.47	7.27	17.15	18.66	21.40	8.53	11.08	11.08	**	+	
	PLACE1004827	3.25	1.26	2.36	5.76	5.15	4.86	3.26	3.82	3.82	**	+	
20	PLACE1004836	2.02	0.78	1.32	3.29	3.51	3.51	1.36	2.69	2.69	**	+	
	PLACE1004838	3.17	2.09	1.89	2.78	2.46	3.36	1.52	3.28	3.28			
	PLACE1004846	1.23	0.56	0.64	2.27	3.76	2.10	1.40	1.24	1.24	*	+	
	PLACE1004842	5.48	1.99	1.07	1.39	1.40	2.34	2.69	3.06	3.06			
	PLACE1004850	3.11	1.83	1.19	2.34	1.99	1.83	2.00	3.44	3.44			
25	PLACE1004868	1.78	1.97	1.38	1.05	1.30	0.94	1.18	1.52	1.52	*	-	
	PLACE1004885	4.12	2.86	3.03	6.17	4.95	6.21	2.81	3.69	3.69	*	+	
	PLACE1004886	1.77	1.59	1.70	1.43	1.55	1.82	2.32	4.30	4.3		*	+
	PLACE1004887	25.24	11.67	14.76	21.81	38.02	28.05	8.65	10.31	10.31			
	PLACE1004896	2.33	1.72	1.45	4.61	4.55	3.16	5.89	7.01	7.01	*	+	**
	PLACE1004900	9.03	4.30	5.53	9.31	10.97	9.80	5.74	6.69	6.69			
30	PLACE1004902	15.98	5.16	8.41	6.64	13.40	8.82	7.56	8.91	8.91			
	PLACE1004904	2.63	1.32	1.15	1.84	2.37	1.90	3.74	3.50	3.5		*	+
	PLACE1004911	1.14	3.11	1.00	4.23	0.30	0.65	0.27	1.36	1.36			
	PLACE1004913	2.14	1.21	1.21	2.7	1.96	3.02	1.97	4.39	4.39			
	PLACE1004918	1.11	0.31	1.10	1.32	1.60	1.48	0.91	1.02	1.02			
35	PLACE1004930	3.51	2.35	1.88	1.71	2.51	2.60	1.12	1.41	1.41			
	PLACE1004934	2.04	1.42	1.26	1.7	2.74	2.49	1.45	1.52	1.52			
	PLACE1004937	5.11	2.46	1.95	3.63	3.54	3.36	2.75	2.15	2.15			
	PLACE1004949	4.03	1.71	2.54	6.88	7.76	8.45	5.04	9.82	9.82	**	+	*
	PLACE1004969	3.48	2.29	1.51	2.73	3.17	3.01	2.31	4.32	4.32			
40	PLACE1004970	0.79	0.82	0.40	0.36	1.00	0.91	0.81	2.69	2.69			
	PLACE1004972	1.78	1.50	1.56	2.23	2.38	3.07	1.16	2.50	2.5	*	+	
	PLACE1004974	3.63	3.03	1.68	3.41	3.31	2.59	1.64	1.70	1.7			
	PLACE1004975	4.46	3.12	2.44	4.13	3.11	5.49	3.51	3.95	3.95			
	PLACE1004979	4.8	5.17	3.63	8.89	10.47	10.51	5.50	6.33	6.33	**	+	*
45	PLACE1004982	12.69	7.06	8.29	13.78	13.06	8.17	7.03	8.87	8.87			
	PLACE1004985	2.12	0.35	0.79	2.05	1.96	1.11	0.99	3.21	3.21			
	PLACE1005003	3.67	1.05	1.88	1.3	2.66	1.79	0.59	2.43	2.43			
	PLACE1005004	1.24	1.06	1.30	1.55	1.31	1.17	1.68	1.83	1.83		**	+
	PLACE1005005	8.08	4.02	3.41	8.61	8.51	8.54	5.01	5.29	5.29			
	PLACE1005011	2.2	1.69	2.79	3	3.06	5.33	3.11	2.57	2.57			
50	PLACE1005026	2.34	1.90	2.06	1.86	2.01	2.93	1.82	1.53	1.53		*	-
	PLACE1005027	4.99	3.43	4.26	7.85	11.24	9.53	3.37	5.57	5.57	**	+	
	PLACE1005031	6.43	2.62	2.97	5.45	4.09	3.20	3.04	3.84	3.84			
	PLACE1005036	7.51	3.86	5.10	9.33	12.02	7.99	3.66	4.98	4.98			
	PLACE1005041	0.87	0.69	0.58	1.76	1.87	1.43	1.58	1.91	1.91	**	+	**
55	PLACE1005046	7.09	3.32	3.54	11.1	10.13	10.18	4.94	5.99	5.99	**	+	
	PLACE1005047	3.57	1.97	1.47	3.2	3.87	3.39	2.49	3.04	3.04			
	PLACE1005052	4.36	2.90	3.32	3.11	2.49	4.07	4.21	4.75	4.75			

Table 311

5	PLACE1004441	3.25	1.90	2.33	4.32	4.15	5.16	3.84	4.52	4.52	*	+	*	+
	PLACE1004446	1.76	2.09	0.72	1.34	1.42	1.87	2.28	2.32	2.32				
	PLACE1004450	0.76	0.23	0.38	0.96	1.30	0.99	0.73	0.72	0.72	*	+		
	PLACE1004451	2.04	1.05	0.94	1.87	2.71	1.33	1.83	2.40	2.4				
	PLACE1004456	13.14	7.90	8.58	15.19	13.06	9.85	9.75	13.11	13.11				
10	PLACE1004458	1.13	0.48	0.38	2.8	2.09	3.55	9.05	9.62	9.62	*	+	**	+
	PLACE1004460	1.24	0.45	0.57	1.15	1.35	1.69	1.34	1.71	1.71		*	+	
	PLACE1004467	6.23	3.77	6.46	8.7	9.58	9.65	5.25	4.76	4.76	*	+		
	PLACE1004471	7.06	5.28	5.80	10.51	12.81	16.26	6.17	7.08	7.08	*	+		
	PLACE1004473	1.57	1.48	1.06	1.91	1.92	2.41	1.84	1.43	1.43	*	+		
	PLACE1004475	17.9	8.89	9.13	27.5	24.29	13.71	28.08	20.33	20.33		*	+	
15	PLACE1004482	2.18	1.39	1.16	1.98	2.90	3.51	2.75	3.78	3.78		*	+	
	PLACE1004491	0.74	0.46	0.72	0.47	1.01	0.56	0.69	1.94	1.94				
	PLACE1004492	33.34	16.09	17.54	17.67	22.65	21.39	20.85	24.45	24.45				
	PLACE1004506	5.1	3.77	3.89	3.53	5.30	4.79	5.63	7.41	7.41		*	+	
	PLACE1004507	2.94	1.98	2.25	1.75	2.11	1.80	2.62	3.67	3.67				
20	PLACE1004510	2.01	2.57	2.33	4.62	4.58	4.58	3.18	2.57	2.57	**	+		
	PLACE1004516	1.04	0.43	0.32	0.6	0.82	1.51	0.69	1.14	1.14				
	PLACE1004518	5.88	3.35	1.73	3.03	3.63	1.95	4.27	3.46	3.46				
	PLACE1004519	3.55	1.36	2.17	1.53	2.33	1.77	1.26	1.42	1.42				
	PLACE1004520	4.8	1.73	3.29	3.58	4.49	2.98	3.20	4.60	4.6				
25	PLACE1004530	7.81	5.59	5.82	2.93	4.17	2.72	3.17	3.36	3.36	*	-	*	-
	PLACE1004545	0.98	1.24	0.71	1.02	1.35	1.28	1.23	1.48	1.48				
	PLACE1004547	3.48	2.58	2.62	3.89	3.59	4.14	3.27	6.00	6	*	+		
	PLACE1004548	5.32	3.02	2.13	5.34	7.57	7.29	2.74	4.90	4.9				
	PLACE1004550	4.75	3.89	2.55	4.32	5.77	4.11	3.73	5.54	5.54				
	PLACE1004551	2.21	1.18	1.01	2.32	3.16	1.67	1.47	1.73	1.73				
30	PLACE1004559	1.69	0.68	1.41	2.2	2.41	1.95	1.58	1.77	1.77	*	+		
	PLACE1004562	7.92	4.63	4.61	12.8	13.69	12.24	11.70	16.91	16.91	**	+	*	+
	PLACE1004564	5.08	3.48	2.94	3.43	4.16	2.75	2.50	3.03	3.03				
	PLACE1004604	1.61	1.65	0.87	1.96	1.66	1.23	6.31	2.27	2.27				
	PLACE1004611	6.51	4.71	3.22	13.38	14.72	11.15	6.91	6.89	6.89	**	+		
35	PLACE1004629	3.8	3.23	3.16	7.62	7.80	6.85	5.92	7.19	7.19	**	+	**	+
	PLACE1004630	4.43	7.59	4.92	4.3	3.84	5.63	3.88	4.82	4.82				
	PLACE1004637	9.71	8.66	5.16	8.97	5.26	6.98	6.87	7.85	7.85				
	PLACE1004645	34.24	15.91	17.01	26.16	30.73	32.52	15.81	17.34	17.34				
	PLACE1004646	3.38	1.74	3.32	3.28	4.81	3.28	2.79	2.82	2.82				
40	PLACE1004648	14.4	8.71	8.36	10.69	11.92	11.82	11.67	15.16	15.16				
	PLACE1004655	41.73	23.86	25.42	40	42.96	45.63	19.14	24.74	24.74				
	PLACE1004658	4.07	3.17	2.80	4.22	4.91	5.38	4.38	3.84	3.84	*	+		
	PLACE1004664	2.14	1.15	0.86	2.2	2.05	3.93	1.74	1.79	1.79				
	PLACE1004672	11.36	7.67	9.44	13.22	15.37	20.21	6.56	12.23	12.23	*	+		
	PLACE1004674	6.89	4.27	3.73	8.23	11.59	6.63	7.24	9.33	9.33		*	+	
45	PLACE1004681	5.36	2.49	2.37	3.93	6.34	2.28	3.03	2.81	2.81				
	PLACE1004686	4.25	1.52	2.69	8.28	8.25	8.23	3.83	5.37	5.37	**	+		
	PLACE1004690	27.35	18.33	25.68	19.28	26.77	23.31	7.55	15.04	15.04		*	-	
	PLACE1004691	4.78	2.55	2.69	4.7	7.55	6.34	2.68	5.61	5.61				
	PLACE1004693	3.07	1.09	1.84	2.44	2.98	3.35	2.53	3.19	3.19				
50	PLACE1004701	23.69	11.94	19.76	25.99	21.50	33.00	23.49	24.31	24.31				
	PLACE1004705	5.61	4.43	3.93	4.87	5.07	5.49	3.83	4.06	4.06				
	PLACE1004708	9.98	7.05	4.96	7.36	17.22	7.73	12.21	12.77	12.77		*	+	
	PLACE1004716	5.47	2.91	3.32	5.79	8.69	5.11	4.23	4.07	4.07				
	PLACE1004722	1.53	1.35	1.55	1.7	3.73	3.67	0.90	2.30	2.3				
55	PLACE1004736	16.73	9.74	14.43	13.11	17.99	18.80	11.66	17.15	17.15				
	PLACE1004737	2.18	1.67	1.79	1.43	3.14	3.32	1.37	1.80	1.8				
	PLACE1004740	6.4	3.17	4.45	6.16	4.57	7.19	5.04	5.92	5.92				

Table 310

5	PLACE1004104	2.15	1.27	0.85	1.43	1.39	2.13	1.09	2.01	2.01				
	PLACE1004113	4.08	1.68	3.31	4.6	4.46	4.54	3.36	3.05	3.05				
	PLACE1004114	2.54	0.84	0.51	1.58	2.53	1.82	2.42	1.88	1.88				
	PLACE1004118	1.98	1.29	1.42	1.63	4.01	2.38	1.61	2.11	2.11				
	PLACE1004128	12.83	9.07	9.04	8.02	8.50	9.63	5.06	6.17	6.17		*	-	
	PLACE1004130	2.24	2.05	1.32	1.83	3.44	3.33	2.12	1.72	1.72				
10	PLACE1004149	18	9.56	12.62	22.09	23.13	25.79	15.85	17.31	17.31	*	+		
	PLACE1004156	8.66	4.78	4.97	11.23	13.14	12.83	5.87	8.14	8.14	*	+		
	PLACE1004160	31.97	23.56	27.55	20.37	16.69	25.95	28.83	35.50	35.5				
	PLACE1004161	12.19	6.98	6.65	7.81	8.30	9.68	8.49	8.65	8.65				
	PLACE1004166	10.59	4.49	3.61	8.56	19.40	8.04	5.20	7.58	7.58				
15	PLACE1004168	9.22	3.40	4.94	7.74	9.05	6.39	5.52	5.88	5.88				
	PLACE1004170	0.56	0.65	1.17	2.02	1.70	2.28	1.72	2.24	2.24	**	+	**	+
	PLACE1004178	5.68	2.50	3.59	4.97	6.58	6.01	4.61	7.20	7.2				
	PLACE1004183	4.44	2.26	4.45	5.52	5.64	5.63	4.08	3.85	3.85				
	PLACE1004197	1.06	1.17	1.74	1.07	1.49	1.13	2.10	1.67	1.67				
20	PLACE1004199	9.96	6.47	8.63	4.5	6.39	5.99	10.80	9.20	9.2				
	PLACE1004203	6.09	3.61	5.37	4.74	4.70	4.68	5.77	5.62	5.62				
	PLACE1004242	7.53	2.60	2.25	8.1	9.90	6.46	4.60	5.49	5.49				
	PLACE1004249	25.51	14.54	13.20	20.9	26.96	19.21	17.71	21.13	21.13				
	PLACE1004255	1.02	0.75	0.36	0.86	1.57	1.36	0.69	1.07	1.07				
25	PLACE1004256	4.42	1.01	3.09	9.24	13.36	13.94	12.44	10.96	10.96	**	+	**	+
	PLACE1004257	4.54	1.21	1.79	4.96	4.55	4.58	3.59	4.84	4.84				
	PLACE1004258	3.59	2.38	2.35	2.98	2.70	2.85	3.20	2.02	2.02				
	PLACE1004270	3.93	3.24	3.36	3.85	4.28	6.05	3.70	3.05	3.05				
	PLACE1004272	4.04	2.85	3.28	3.85	5.74	5.17	3.42	6.23	6.23				
	PLACE1004273	83.7	57.27	49.34	101.5	84.19	78.07	49.24	46.63	46.63				
30	PLACE1004274	2.95	0.92	1.52	1.53	2.26	1.62	1.54	1.70	1.7				
	PLACE1004277	4.89	3.63	3.77	5.98	6.33	5.84	3.49	5.35	5.35	*	+		
	PLACE1004279	4.14	2.37	2.56	4.12	4.89	5.01	2.41	5.41	5.41				
	PLACE1004282	4.87	1.71	2.16	3.7	2.78	3.26	3.33	4.30	4.3				
	PLACE1004284	5.6	3.43	5.55	7.94	7.12	9.08	5.18	6.08	6.08	*	+		
35	PLACE1004289	4.45	2.76	2.32	4.87	4.64	6.03	3.57	3.74	3.74				
	PLACE1004299	3.82	1.87	1.73	3.07	2.88	4.42	3.05	2.95	2.95				
	PLACE1004302	2.2	0.86	0.90	1.74	3.32	2.03	1.19	1.35	1.35				
	PLACE1004305	3.85	2.26	1.59	1.85	1.24	2.43	2.28	2.58	2.58				
	PLACE1004316	5.43	2.71	3.07	1.96	3.30	2.21	2.72	4.32	4.32				
40	PLACE1004322	1.43	0.69	0.73	1.49	2.28	1.46	1.11	2.06	2.06				
	PLACE1004325	13.88	6.16	7.35	9.82	9.01	12.35	11.00	10.37	10.37				
	PLACE1004332	3.01	1.40	1.75	1.66	1.82	2.98	2.54	3.00	3				
	PLACE1004336	9.91	5.69	5.62	10.43	10.12	10.42	6.74	8.77	8.77				
	PLACE1004346	3.07	2.03	1.73	2.75	2.78	2.82	1.63	2.50	2.5				
	PLACE1004358	17.58	10.51	10.45	12.26	12.55	14.79	12.38	16.11	16.11				
45	PLACE1004376	21.68	10.31	10.00	13.3	12.08	12.35	12.00	16.69	16.69				
	PLACE1004384	3.8	1.61	2.13	4.74	4.70	5.37	3.12	3.81	3.81	*	+		
	PLACE1004385	1.9	0.86	0.50	0.57	1.48	1.53	0.60	1.25	1.25				
	PLACE1004388	3.6	1.83	1.85	3.69	4.57	4.12	1.57	1.95	1.95				
	PLACE1004405	0.61	1.07	0.82	0.21	0.91	1.17	2.14	2.86	2.86		**	+	
50	PLACE1004407	5.17	3.33	4.50	6.41	4.17	7.01	4.80	3.58	3.58				
	PLACE1004424	1.66	0.59	0.46	0.66	0.44	2.14	0.23	0.46	0.46				
	PLACE1004425	1.47	0.52	1.48	2.94	3.61	2.12	1.72	2.14	2.14	*	+		
	PLACE1004427	2.86	1.31	1.07	1.87	2.05	1.81	1.96	3.44	3.44				
	PLACE1004428	3.96	2.20	1.76	4.03	5.57	4.86	3.58	3.24	3.24				
	PLACE1004433	6.32	3.82	4.97	5.63	4.68	5.19	2.95	4.99	4.99				
55	PLACE1004435	7.56	3.49	4.09	10.74	10.16	12.36	5.74	11.13	11.13	*	+		
	PLACE1004437	7.97	3.59	4.68	4.42	7.20	5.02	5.17	3.07	3.07				

Table 309

5	PLACE1003660	3.6	2.90	2.17	3.69	3.98	5.22	2.65	3.15	3.15				
	PLACE1003669	3.72	1.83	1.76	4.6	5.24	5.00	3.90	4.38	4.38	*	+		
	PLACE1003670	15.52	7.07	8.39	9.52	9.26	10.68	8.82	8.03	8.03				
	PLACE1003671	4.94	3.13	2.14	3.75	4.23	3.08	3.20	4.09	4.09				
	PLACE1003697	3.08	0.80	1.06	3.54	2.83	2.50	7.26	8.03	8.03		**	+	
10	PLACE1003704	11.2	5.78	7.63	14.43	11.92	13.54	6.97	9.55	9.55	*	+		
	PLACE1003709	4.98	0.98	1.82	0.79	0.50	1.26	1.00	1.96	1.96				
	PLACE1003711	5.06	3.03	2.94	3.49	4.07	3.66	3.26	4.30	4.3				
	PLACE1003723	4.06	2.93	3.32	6.92	5.34	6.03	4.19	5.65	5.65	**	+	*	+
	PLACE1003724	9.61	5.81	6.68	10.85	14.36	13.13	7.86	7.40	7.4	*	+		
	PLACE1003737	1.82	0.70	1.20	1.4	2.78	1.47	0.99	1.14	1.14				
15	PLACE1003738	4.42	2.23	2.32	2.25	3.92	3.77	2.75	4.94	4.94				
	PLACE1003742	4.22	2.78	3.39	5.61	5.88	6.94	5.65	8.11	8.11	*	+	*	+
	PLACE1003744	10.38	5.06	4.96	6	6.16	5.58	7.58	7.15	7.15				
	PLACE1003758	2.34	1.24	1.52	3.36	2.67	2.23	1.96	3.95	3.95				
	PLACE1003760	12.25	10.24	12.40	34.22	35.40	36.07	24.12	29.73	29.73	**	+	**	+
20	PLACE1003762	3.15	2.22	1.75	4.15	5.03	5.81	2.19	3.25	3.25	*	+		
	PLACE1003765	3.6	2.58	2.17	4.49	5.32	6.00	3.44	2.48	2.48	*	+		
	PLACE1003768	2.32	0.82	0.97	3.88	3.45	2.85	1.41	2.13	2.13	*	+		
	PLACE1003771	1.14	0.42	0.47	3.82	4.60	4.57	2.76	2.88	2.88	**	+	**	+
	PLACE1003772	15.91	10.99	11.28	22.4	31.67	17.46	9.36	14.35	14.35				
25	PLACE1003783	1.42	1.64	0.56	2.3	1.57	1.94	2.32	2.86	2.86		*	+	
	PLACE1003784	1.03	0.77	0.68	0.97	1.55	1.05	1.26	0.82	0.82				
	PLACE1003788	1.09	0.76	0.74	1.58	0.81	1.20	1.20	1.12	1.12				
	PLACE1003795	3.57	3.15	3.29	4.82	6.11	5.73	4.14	3.97	3.97	**	+	**	+
	PLACE1003827	4.25	3.25	4.26	3.97	4.73	4.26	4.86	4.32	4.32				
30	PLACE1003833	5.49	4.93	3.72	7.29	6.79	7.39	4.43	6.36	6.36	*	+		
	PLACE1003839	15.63	9.41	9.25	19.2	21.48	17.62	11.21	10.43	10.43	*	+		
	PLACE1003845	7.01	4.24	4.12	7.35	7.87	5.86	10.74	9.90	9.9		**	+	
	PLACE1003850	8.77	5.05	5.31	6.16	11.18	6.64	4.92	6.94	6.94				
	PLACE1003852	1.98	0.95	1.19	2.52	2.43	1.55	2.10	2.14	2.14				
	PLACE1003858	1.86	1.56	1.42	0.9	3.73	1.64	1.18	2.61	2.61				
35	PLACE1003861	3.4	2.90	2.88	4.73	4.58	4.45	3.62	4.50	4.5	**	+	*	+
	PLACE1003864	2.18	1.73	1.70	2.15	2.33	2.94	1.58	1.90	1.9				
	PLACE1003878	6.85	4.56	2.90	9.94	13.82	9.81	3.57	5.78	5.78	*	+		
	PLACE1003885	3.97	2.09	1.62	4.09	4.19	2.32	1.33	1.78	1.78				
	PLACE1003886	6.25	3.53	4.72	4.17	5.68	4.34	4.84	5.28	5.28				
40	PLACE1003888	2.5	2.14	1.29	2.33	2.24	2.51	1.57	1.20	1.2				
	PLACE1003892	0.63	0.82	0.35	1.2	1.75	1.76	1.12	1.37	1.37	*	+	*	+
	PLACE1003900	2.12	3.11	2.67	2.84	3.42	2.21	3.08	3.08	3.08				
	PLACE1003902	2.67	2.85	2.44	2.17	3.06	3.38	2.09	2.93	2.93				
	PLACE1003903	3.07	2.52	2.90	2.6	2.59	4.30	2.16	2.90	2.9				
45	PLACE1003915	2.93	1.59	2.90	5.14	5.88	4.52	4.31	3.51	3.51	*	+		
	PLACE1003918	6.79	4.23	4.22	10.17	14.99	6.89	4.36	4.29	4.29				
	PLACE1003923	2.38	0.83	1.07	2.53	2.23	1.52	2.50	2.86	2.86				
	PLACE1003932	6.11	3.09	2.41	4.35	4.43	3.41	2.40	3.60	3.6				
	PLACE1003936	3.26	3.57	2.07	4.2	6.78	4.35	2.70	3.36	3.36				
	PLACE1003966	2.8	1.71	1.81	3.31	3.64	3.17	2.63	3.00	3	*	+		
50	PLACE1003968	3.23	5.15	4.73	6.02	6.25	7.79	6.80	6.11	6.11	*	+	*	+
	PLACE1004018	3.13	3.22	4.15	3.49	4.47	4.76	3.66	5.03	5.03				
	PLACE1004020	8.8	4.07	3.96	10.35	11.19	10.96	7.35	8.57	8.57	*	+		
	PLACE1004028	2.58	0.63	1.65	1.41	1.48	2.22	1.55	1.12	1.12				
	PLACE1004034	14.58	6.23	5.10	4.38	6.92	4.97	2.01	1.93	1.93				
55	PLACE1004042	13.64	6.23	8.71	10.86	12.72	14.34	11.39	17.16	17.16				
	PLACE1004078	4.38	2.37	2.45	5.69	4.75	6.84	3.40	5.09	5.09	*	+		
	PLACE1004103	7.95	4.34	4.17	15.49	14.70	18.99	9.99	10.73	10.73	**	+	*	+

Table 308

5	PLACE1003372	4.86	3.69	3.10	6.36	6.08	6.40	5.24	6.26	6.26	*	+	*	+
	PLACE1003373	4.59	2.14	-1.77	6.44	8.87	7.14	3.34	3.58	3.58	*	+		
	PLACE1003375	1.64	2.20	2.31	1.72	2.46	2.62	1.19	1.43	1.43			*	-
	PLACE1003378	2.12	1.60	1.04	2.23	1.84	1.68	2.18	2.69	2.69				
	PLACE1003383	2.45	1.53	0.51	2.22	2.04	0.76	1.14	1.36	1.36				
	PLACE1003394	8.16	3.88	4.89	10.77	12.17	8.54	8.17	10.02	10.02	*	+		
10	PLACE1003401	3.67	0.79	0.99	1.2	1.46	1.82	0.45	1.86	1.86				
	PLACE1003405	6.01	6.00	6.98	4.76	7.61	8.04	6.47	7.65	7.65				
	PLACE1003407	4.49	4.04	3.71	5.05	5.22	5.15	5.12	5.61	5.61	*	+	**	+
	PLACE1003420	4.75	4.07	3.59	7.55	10.89	8.12	4.15	6.01	6.01	*	+		
	PLACE1003428	2.19	2.41	3.05	3.29	4.02	4.47	2.16	2.43	2.43	*	+		
15	PLACE1003432	7.17	3.85	3.68	4.37	7.22	7.66	3.81	6.34	6.34				
	PLACE1003438	9.06	3.37	4.39	5.86	7.12	5.43	5.87	7.15	7.15				
	PLACE1003452	3.13	1.08	2.21	1.29	5.01	2.29	2.22	2.52	2.52				
	PLACE1003454	8.4	4.68	5.18	7.33	6.34	9.17	4.92	7.46	7.46				
	PLACE1003455	13.75	5.01	6.05	6.83	8.91	9.83	8.45	9.21	9.21				
20	PLACE1003456	7.28	4.38	4.13	10.64	12.00	13.60	7.62	7.20	7.2	**	+		
	PLACE1003460	7.84	3.76	6.10	10.15	7.44	7.77	6.55	7.66	7.66				
	PLACE1003478	3.33	0.56	0.93	2.01	1.78	1.24	0.65	0.96	0.96				
	PLACE1003484	7.55	4.57	2.88	11.32	16.35	7.83	7.21	9.47	9.47				
	PLACE1003493	14.03	6.96	6.73	11.22	11.97	14.63	9.74	9.34	9.34				
25	PLACE1003583	42.11	19.93	34.28	29.63	36.26	35.89	25.50	29.49	29.49				
	PLACE1003585	2.24	1.06	0.89	0.91	0.90	1.59	2.08	1.73	1.73				
	PLACE1003516	1.01	0.49	0.89	2.17	2.40	2.58	1.68	1.86	1.86	**	+	**	+
	PLACE1003519	39.78	23.99	30.04	55.6	50.01	57.71	22.97	28.09	28.09	*	+		
	PLACE1003520	45.85	22.30	34.27	66.52	30.94	72.87	38.79	44.73	44.73				
	PLACE1003521	1.43	0.65	0.89	2.33	3.32	0.95	2.10	3.87	3.87		*	+	
30	PLACE1003525	15.69	8.19	8.09	12.57	19.45	12.58	15.38	18.26	18.26				
	PLACE1003528	126.72	75.71	77.51	102.34	128.72	89.84	56.09	57.39	57.39				
	PLACE1003529	10.31	6.25	7.90	10.63	11.63	11.54	9.31	9.78	9.78				
	PLACE1003537	3.45	1.76	2.18	3.36	4.60	3.48	5.58	5.15	5.15		**	+	
	PLACE1003549	3.96	2.80	3.67	4.57	2.88	5.08	2.97	4.32	4.32				
35	PLACE1003553	6.15	2.35	3.07	4.85	4.12	5.00	3.14	3.29	3.29				
	PLACE1003566	5.25	2.36	2.80	5.45	5.03	6.90	4.92	5.27	5.27				
	PLACE1003568	1.39	1.43	0.56	1.66	1.56	1.27	1.01	0.83	0.83				
	PLACE1003573	2.04	1.89	1.09	2.09	2.81	1.71	1.61	1.69	1.69				
	PLACE1003575	3.94	2.36	1.55	4.2	5.03	5.48	3.67	2.41	2.41	*	+		
40	PLACE1003583	1.25	0.21	0.91	0.63	1.54	1.28	1.19	0.85	0.85				
	PLACE1003584	3.17	2.52	1.33	5.76	4.75	5.94	2.30	3.30	3.3	**	+		
	PLACE1003592	6.37	4.34	3.44	8.54	12.20	11.57	7.98	8.85	8.85	*	+	*	+
	PLACE1003593	0.73	1.09	0.64	1.3	1.69	1.81	0.49	1.57	1.57	*	+		
	PLACE1003594	16.13	4.42	11.69	14.87	17.87	21.56	10.51	11.29	11.29				
45	PLACE1003596	5.64	5.18	5.93	10.49	15.28	7.57	7.20	9.60	9.6		*	+	
	PLACE1003598	13.48	8.08	6.25	8.41	8.96	8.69	7.81	8.78	8.78				
	PLACE1003602	3.72	2.13	1.45	3.5	3.57	3.37	2.64	3.45	3.45				
	PLACE1003605	18.39	10.93	10.02	16.96	17.66	21.30	9.74	14.50	14.5				
	PLACE1003611	3.07	0.86	1.19	2.62	2.97	3.49	1.69	2.05	2.05				
	PLACE1003618	2.42	0.71	0.96	1.64	1.41	1.56	1.78	2.12	2.12				
50	PLACE1003625	3.62	1.30	2.39	3.11	4.04	4.15	3.30	3.49	3.49				
	PLACE1003626	13.07	5.94	8.16	14.48	13.10	14.74	12.62	11.51	11.51				
	PLACE1003630	3.48	2.42	1.94	3.18	3.39	2.97	3.11	3.27	3.27				
	PLACE1003635	2.04	1.03	1.44	2.07	2.17	2.34	1.81	1.67	1.67				
	PLACE1003638	3.27	2.36	1.79	4.52	4.52	3.82	3.33	3.31	3.31	*	+		
55	PLACE1003644	3.31	2.33	2.10	5.21	5.95	5.73	4.05	4.05	4.05	**	+	*	+
	PLACE1003654	4.23	1.54	1.89	1.81	2.78	2.00	0.89	2.32	2.32				
	PLACE1003656	2.23	0.80	1.38	1.4	1.47	1.90	1.48	2.10	2.1				

Table 307

	PLACE1002967	5.1	2.51	3.09	6.76	6.45	5.80	4.18	4.06	4.06	*	+		
5	PLACE1002968	1.23	0.90	0.78	1.96	2.73	1.63	1.67	2.25	2.25	*	+	**	+
	PLACE1002976	14.62	6.59	8.58	14.43	17.61	21.24	10.88	15.26	15.26				
	PLACE1002991	9.09	3.33	5.17	10.69	10.94	9.19	4.59	4.38	4.38				
	PLACE1002993	4.97	3.72	3.40	7.49	6.57	6.94	4.40	4.67	4.67	**	+		
	PLACE1002996	4.17	2.53	2.14	3.73	3.53	2.43	2.53	3.20	3.2				
10	PLACE1003010	14.09	9.21	8.66	11.39	9.01	12.05	11.70	11.85	11.85				
	PLACE1003025	3.37	1.92	1.25	3.12	3.46	2.82	2.56	2.83	2.83				
	PLACE1003027	2.78	1.30	1.63	3.36	4.14	4.94	2.51	3.33	3.33	*	+		
	PLACE1003044	5.29	2.38	3.63	5.05	4.60	4.39	4.30	3.74	3.74				
	PLACE1003045	1.31	0.14	0.41	1.12	0.74	1.58	0.92	1.66	1.66				
15	PLACE1003052	5.81	2.44	2.52	4.24	6.72	5.03	2.74	4.06	4.06				
	PLACE1003083	1.98	0.63	0.30	1.59	1.48	1.45	1.09	1.36	1.36				
	PLACE1003085	8.86	4.56	4.41	4.48	5.13	3.76	5.79	5.25	5.25				
	PLACE1003092	4.95	2.80	2.49	4.61	7.21	5.11	3.15	5.59	5.59				
	PLACE1003097	2.48	1.08	1.75	2.13	2.19	3.46	1.83	1.87	1.87				
20	PLACE1003100	5.55	3.04	3.54	4.48	2.63	4.78	3.66	4.38	4.38				
	PLACE1003108	2.43	2.01	1.88	3.79	4.20	5.56	3.02	3.15	3.15	*	+	**	+
	PLACE1003115	5.59	4.45	4.08	5.2	3.47	4.38	3.94	4.36	4.36				
	PLACE1003120	9.1	5.05	6.99	11.92	11.69	8.39	4.33	5.35	5.35				
	PLACE1003135	7.15	3.42	2.81	2	1.71	2.50	1.33	2.53	2.53				
25	PLACE1003136	9.4	3.19	5.96	7.56	7.72	8.01	6.80	8.18	8.18				
	PLACE1003141	1.43	1.20	0.97	1.12	1.71	2.12	1.29	2.62	2.62				
	PLACE1003145	1.17	1.98	1.88	1.29	0.85	1.19	1.52	2.74	2.74				
	PLACE1003147	3.88	1.84	2.10	3.04	3.09	5.16	2.94	6.44	6.44				
	PLACE1003153	2.04	1.22	1.34	1.76	3.27	2.50	1.12	2.13	2.13				
30	PLACE1003163	5.21	2.54	2.21	3.71	2.70	3.59	1.58	3.29	3.29				
	PLACE1003172	17.21	13.29	11.63	20.51	17.81	16.21	12.82	14.76	14.76				
	PLACE1003174	1.86	0.95	0.96	2.33	2.68	2.13	2.07	2.85	2.85	*	+	*	+
	PLACE1003176	1.87	0.85	0.99	0.69	1.79	1.46	1.77	2.02	2.02				
	PLACE1003181	2.42	1.29	1.30	1.36	1.88	1.93	2.33	2.76	2.76				
	PLACE1003184	4.02	2.35	1.57	1.09	1.42	1.68	2.02	2.95	2.95				
35	PLACE1003190	12.59	7.17	8.42	3.7	4.03	4.95	5.55	3.22	3.22	*	-	*	-
	PLACE1003200	0.16	0.08	0.11	0.98	0.55	0.76	0.91	1.63	1.63	**	+	**	+
	PLACE1003205	10.63	4.75	4.99	13.42	19.02	15.56	5.60	9.62	9.62	*	+		
	PLACE1003209	1.33	0.58	0.91	1.06	1.71	1.13	1.44	1.84	1.84		*	+	
	PLACE1003214	3.74	1.92	0.96	2.48	3.08	2.07	2.80	1.58	1.58				
40	PLACE1003229	4.01	2.47	1.89	4.67	6.17	5.71	3.46	3.20	3.2	*	+		
	PLACE1003238	0.55	1.29	0.72	1.01	1.04	1.42	1.89	4.82	4.82		*	+	
	PLACE1003249	4.21	2.68	2.29	5.89	6.34	7.49	3.21	4.18	4.18	**	+		
	PLACE1003256	15.42	10.76	11.86	18.06	20.59	21.48	20.54	17.58	17.58	*	+	*	+
	PLACE1003258	1.59	3.70	0.75	1.91	1.78	1.15	1.24	1.39	1.39				
45	PLACE1003279	5.6	4.25	1.88	7.33	8.87	7.26	3.36	5.26	5.26	*	+		
	PLACE1003294	5.96	3.04	2.55	5.19	4.93	5.17	2.65	4.69	4.69				
	PLACE1003296	3.69	1.73	1.93	4.06	3.41	2.82	2.94	3.29	3.29				
	PLACE1003297	6.38	2.82	3.60	6.92	8.35	6.63	3.36	5.38	5.38				
	PLACE1003302	6.92	3.76	5.11	9.34	12.52	9.10	7.08	7.90	7.9	*	+		
	PLACE1003334	0.67	1.10	1.68	2.93	3.55	4.00	2.24	3.22	3.22	**	+	*	+
50	PLACE1003337	10.11	6.39	4.50	7.53	11.58	10.30	6.19	6.17	6.17				
	PLACE1003342	1.8	1.54	1.48	1.86	2.85	3.11	2.71	3.37	3.37		**	+	
	PLACE1003343	0.54	0.36	0.34	0.71	0.79	1.22	0.47	0.55	0.55	*	+		
	PLACE1003344	24.27	18.53	13.01	17.74	21.43	22.39	17.02	18.57	18.57				
	PLACE1003353	17.73	10.09	9.18	16.17	17.86	14.13	8.53	9.79	9.79				
55	PLACE1003361	5.88	2.41	3.54	11.34	11.94	9.89	3.99	5.89	5.89	**	+		
	PLACE1003366	6.48	3.29	5.05	6.5	8.30	6.96	4.61	4.25	4.25				
	PLACE1003369	2.89	2.16	1.46	3.79	2.79	2.98	2.58	2.98	2.98				

Table 306

5	PLACE1002499	3.87	1.99	3.01	5.9	5.94	5.28	3.14	5.21	5.21	**	+		
	PLACE1002500	3.82	3.46	3.57	5.63	5.50	7.08	4.28	4.54	4.54	**	+	**	+
	PLACE1002514	2.68	2.18	1.93	2.67	2.24	2.48	3.81	2.98	2.98			*	+
	PLACE1002518	3.35	3.89	3.09	9.93	9.60	8.45	4.26	3.29	3.29	**	+		
	PLACE1002529	1.4	1.36	1.04	1.77	2.14	1.22	1.26	1.64	1.64				
	PLACE1002532	8.72	6.46	7.19	6.81	6.68	6.18	7.67	10.12	10.12				
10	PLACE1002536	4.9	1.91	3.90	5.09	4.55	3.56	4.82	3.96	3.96				
	PLACE1002537	3.14	1.37	1.42	3.63	3.37	4.11	2.67	3.94	3.94	*	+		
	PLACE1002539	3.39	2.92	3.22	4.41	4.54	5.47	3.82	4.68	4.68	*	+	*	+
	PLACE1002547	5.53	5.37	5.59	8.39	7.22	9.28	8.89	10.18	10.18	**	+	**	+
	PLACE1002571	4.43	2.94	4.05	4.84	4.88	7.44	3.32	5.08	5.08				
15	PLACE1002578	5.19	3.96	3.76	12.25	10.98	12.86	5.35	7.25	7.25	**	+	*	+
	PLACE1002583	1.66	0.32	1.44	1.04	1.08	1.16	1.18	0.97	0.97				
	PLACE1002591	3.86	2.09	2.10	2.84	2.83	2.65	2.44	2.62	2.62				
	PLACE1002598	3.84	2.11	2.49	1.35	1.31	2.14	2.05	2.70	2.7				
	PLACE1002604	2.65	2.17	1.64	2.8	3.94	3.24	2.45	2.54	2.54				
20	PLACE1002612	8.01	6.63	6.63	12.1	11.80	12.23	8.71	11.33	11.33	**	+	*	+
	PLACE1002625	2.58	1.69	1.51	2.59	2.61	4.00	1.54	3.25	3.25				
	PLACE1002638	2.18	2.76	3.22	4.42	3.44	3.29	2.42	3.06	3.06				
	PLACE1002655	3.25	4.16	4.18	10.46	6.84	7.33	3.29	5.31	5.31	*	+		
	PLACE1002665	4.13	3.33	2.98	6.38	8.85	5.64	3.53	3.33	3.33	*	+		
	PLACE1002685	5.53	3.42	2.72	4.03	3.03	3.10	2.10	4.59	4.59				
25	PLACE1002692	8.81	6.44	4.56	11.47	12.80	10.48	5.21	5.98	5.98	*	+		
	PLACE1002714	6.78	4.06	3.36	6.88	8.05	6.09	4.05	4.56	4.56				
	PLACE1002721	6.84	4.40	5.49	7.7	6.58	8.37	3.72	4.94	4.94				
	PLACE1002722	0.74	0.78	0.84	1.77	1.11	1.67	0.84	1.69	1.69	*	+		
	PLACE1002726	3.49	5.71	5.81	8.46	6.47	7.89	5.02	5.08	5.08				
30	PLACE1002756	3.26	2.58	3.14	6.13	6.45	6.35	3.62	4.77	4.77	**	+	*	+
	PLACE1002768	3.97	1.25	1.67	3.3	2.50	2.51	2.78	3.00	3				
	PLACE1002772	1.35	0.09	0.96	0.92	1.25	1.40	1.29	1.37	1.37				
	PLACE1002775	14.42	7.79	9.64	10.55	11.21	17.27	12.36	11.82	11.82				
	PLACE1002780	1.98	1.39	1.23	1.94	2.57	3.18	3.07	5.79	5.79			*	+
35	PLACE1002782	3.02	0.85	1.61	1.99	1.37	3.05	1.65	1.52	1.52				
	PLACE1002794	2.49	1.48	2.20	1.75	2.76	3.63	2.18	2.11	2.11				
	PLACE1002795	1.27	0.70	0.60	1.08	1.69	1.49	0.76	0.93	0.93				
	PLACE1002811	3.67	1.25	0.81	2.9	2.86	1.50	2.33	2.50	2.5				
	PLACE1002815	5.44	2.94	2.29	7.27	15.00	10.36	12.84	16.74	16.74	*	+	**	+
40	PLACE1002816	8.2	3.96	3.92	6.25	6.25	6.46	5.01	6.23	6.23				
	PLACE1002822	3.34	1.86	2.08	2.36	4.04	3.46	2.71	3.36	3.36				
	PLACE1002833	7.79	2.79	4.10	8.87	10.15	5.51	5.02	6.82	6.82				
	PLACE1002834	10.13	4.35	5.31	13.91	16.58	13.43	4.69	8.55	8.55	*	+		
	PLACE1002835	10.05	3.42	5.57	6.59	4.99	6.00	6.06	6.48	6.48				
	PLACE1002839	1.69	0.45	1.30	1.57	1.51	2.09	0.58	1.24	1.24				
45	PLACE1002851	0.76	0.42	0.82	8.19	10.42	3.82	2.14	3.56	3.56	*	+	**	+
	PLACE1002853	2.74	1.59	0.90	3.41	5.47	2.91	2.59	4.00	4				
	PLACE1002881	6.4	4.73	3.84	9.33	10.46	8.15	5.45	5.65	5.65	*	+		
	PLACE1002901	24.97	13.25	21.58	24.2	26.27	27.37	19.88	33.42	33.42				
	PLACE1002904	1.92	2.64	1.54	1.71	2.53	1.56	1.90	1.68	1.68				
50	PLACE1002905	3.55	1.81	2.93	4.92	3.38	5.48	3.25	4.03	4.03				
	PLACE1002908	3.03	1.68	2.53	3.06	3.13	3.59	2.41	3.70	3.7				
	PLACE1002911	17.43	10.47	13.64	12.95	9.50	14.36	14.97	14.68	14.68				
	PLACE1002941	4.78	1.82	1.86	3.71	4.48	2.93	2.39	2.15	2.15				
	PLACE1002950	9.17	5.50	4.49	8.77	14.38	6.75	5.85	4.90	4.9				
55	PLACE1002955	30.87	12.83	9.17	24.34	31.40	20.44	28.45	32.62	32.62				
	PLACE1002958	12.6	6.07	8.49	13.92	20.27	16.63	20.63	26.24	26.24	*	+	**	+
	PLACE1002962	1.57	0.87	0.79	1.15	2.51	1.72	1.24	1.43	1.43				

Table 305

5	PLACE1002081	1.99	0.89	1.77	2.72	4.23	2.35	2.10	2.07				
	PLACE1002090	14.44	6.66	9.78	10.42	12.14	11.62	5.32	7.78	7.78			
	PLACE1002095	6.66	3.83	6.14	8.67	7.29	9.40	5.73	7.69	7.69			
	PLACE1002102	11.71	6.09	6.01	11.63	6.93	8.62	6.39	8.11	8.11			
	PLACE1002109	2.46	1.22	1.40	2.6	4.68	2.17	2.82	2.11	2.11			
	PLACE1002115	3.01	0.88	0.58	1.13	2.98	1.33	0.18	1.10	1.1			
10	PLACE1002119	18.69	14.15	17.17	28.94	38.25	31.55	24.25	29.45	29.45	**	+	**
	PLACE1002140	7.37	4.29	6.46	6.39	6.75	7.33	4.91	5.86	5.86			
	PLACE1002150	2.02	1.18	2.19	3.93	4.63	3.78	3.27	2.55	2.55	**	+	
	PLACE1002153	6.36	3.80	4.46	7.01	6.47	4.93	5.54	4.93	4.93			
	PLACE1002157	2.68	1.47	1.39	4.12	3.06	4.68	2.90	3.69	3.69	*	+	*
15	PLACE1002163	7.63	2.62	3.61	7.02	7.14	5.85	5.07	6.08	6.08			
	PLACE1002168	4.33	2.82	2.86	4.8	4.18	3.05	4.14	4.00	4			
	PLACE1002170	2.98	1.54	1.88	1.56	1.84	1.46	1.96	1.92	1.92			
	PLACE1002171	13.45	7.42	8.57	6.89	9.10	5.13	2.02	3.14	3.14		*	-
	PLACE1002180	1.81	0.89	1.51	3.13	3.65	3.26	1.39	2.44	2.44	**	+	
20	PLACE1002184	2.38	1.68	1.24	6.52	7.00	7.36	6.04	5.01	5.01	**	+	**
	PLACE1002200	3.74	3.15	2.61	3.65	2.78	3.93	3.98	4.06	4.06			
	PLACE1002205	1.24	0.51	0.69	2.33	2.64	4.75	1.98	1.74	1.74	*	+	*
	PLACE1002213	8.87	4.30	5.26	10.21	8.63	11.56	6.15	7.84	7.84			
	PLACE1002219	1.89	0.82	0.74	1.44	2.66	1.62	0.97	0.77	0.77			
	PLACE1002227	4.82	2.81	1.66	4.34	4.54	4.85	2.92	3.36	3.36			
25	PLACE1002253	3.86	2.60	1.93	1.41	2.78	1.93	2.88	2.14	2.14			
	PLACE1002256	1.83	0.92	1.11	2.87	3.97	2.85	1.91	3.59	3.59	*	+	*
	PLACE1002259	3.19	1.70	1.57	6.62	7.59	6.60	5.13	4.07	4.07	**	+	*
	PLACE1002285	1.77	0.92	0.70	2.37	1.34	1.10	1.30	2.28	2.28			
	PLACE1002301	3.7	3.54	3.53	4.57	5.90	8.65	6.82	8.88	8.88		**	+
30	PLACE1002310	2.48	1.29	1.37	3.99	3.09	4.29	7.69	9.72	9.72	*	+	**
	PLACE1002311	3.44	2.13	1.55	3.07	3.48	2.34	2.76	2.45	2.45			
	PLACE1002319	4.6	2.18	2.82	2.38	2.25	2.70	1.39	2.13	2.13			
	PLACE1002329	4.19	2.99	2.11	3.47	3.41	5.33	3.47	4.66	4.66			
	PLACE1002333	1.41	1.34	1.43	2.55	1.71	1.03	1.08	1.25	1.25		*	-
35	PLACE1002342	3.55	2.39	2.93	7.53	5.67	7.31	3.57	4.06	4.06	**	+	
	PLACE1002343	3.11	2.65	3.16	2.86	3.12	2.88	2.90	5.44	5.44			
	PLACE1002355	3.89	1.69	1.70	3.76	3.03	3.60	3.29	2.58	2.58			
	PLACE1002358	3.55	2.39	2.49	3.8	3.99	2.81	2.23	2.70	2.7			
	PLACE1002359	8	4.42	4.71	3.91	5.64	5.32	4.07	5.01	5.01			
40	PLACE1002374	14.74	8.20	8.86	9.64	10.72	8.98	11.09	14.20	14.2			
	PLACE1002376	7.57	5.16	5.69	9.15	8.50	11.00	8.02	8.55	8.55	*	+	*
	PLACE1002379	3.61	3.25	3.56	3.36	3.66	3.11	4.20	4.20	4.2		**	+
	PLACE1002386	5.82	2.32	2.77	4.29	2.48	5.32	6.23	7.32	7.32		*	+
	PLACE1002395	5.61	3.00	2.85	4.9	4.34	4.62	4.54	4.04	4.04			
	PLACE1002399	2.61	1.20	1.56	3.06	2.87	4.76	3.56	3.31	3.31		*	+
45	PLACE1002407	4.59	2.71	2.96	2.81	2.75	3.28	1.95	2.26	2.26			
	PLACE1002433	5.13	3.15	3.02	4.68	5.35	5.03	2.23	2.89	2.89			
	PLACE1002437	3.54	1.57	2.70	3.76	3.24	3.17	2.59	4.55	4.55			
	PLACE1002438	1.21	1.24	1.22	1.63	1.79	2.23	2.23	2.71	2.71	*	+	**
	PLACE1002446	5.14	2.19	2.50	4.51	3.25	4.13	6.69	7.98	7.98		*	+
50	PLACE1002447	2.92	2.41	2.19	1.36	3.06	1.99	2.41	2.93	2.93			
	PLACE1002450	1.44	2.03	1.72	3.08	3.13	3.49	2.56	2.00	2	**	+	
	PLACE1002462	2.28	1.70	1.59	1.95	1.67	3.14	1.58	2.39	2.39			
	PLACE1002465	3.1	2.98	2.42	2.1	3.45	3.05	2.13	2.02	2.02		*	-
	PLACE1002474	2.91	2.82	2.76	8.43	9.88	7.40	6.02	7.81	7.81	**	+	**
55	PLACE1002477	8.13	3.74	5.00	11.28	9.50	9.10	9.42	12.59	12.59	*	+	*
	PLACE1002493	1.9	2.19	1.21	1.77	1.63	1.93	2.11	3.26	3.26			
	PLACE1002497	2.74	1.52	2.43	1.73	1.95	2.51	2.01	3.14	3.14			

Table 304

	PLACE1001728	1.5	1.02	0.69	1.1	0.60	1.41	1.40	1.39	1.39				
5	PLACE1001729	6.79	3.57	3.61	3.84	3.10	4.27	2.54	6.08	6.08				
	PLACE1001739	9.94	5.41	6.00	8.04	5.84	6.73	6.37	6.11	6.11				
	PLACE1001740	1.57	0.32	0.49	0.97	1.11	1.42	1.06	0.82	0.82				
	PLACE1001745	5.8	3.72	3.68	4.06	4.53	4.47	4.22	4.88	4.88				
	PLACE1001746	3.57	1.52	1.71	4.99	5.18	6.01	3.66	5.62	5.62	*	+	*	+
10	PLACE1001748	4.5	2.90	2.37	5.53	4.76	3.57	3.80	5.19	5.19				
	PLACE1001753	3.51	2.28	3.04	2.88	3.35	3.77	3.11	5.17	5.17				
	PLACE1001756	12.16	6.46	7.86	8.59	7.90	8.09	4.55	8.41	8.41				
	PLACE1001760	8.72	4.93	5.18	11.47	11.77	9.41	7.48	10.20	10.2	*	+		
	PLACE1001767	6.27	4.18	2.75	5.86	5.81	6.64	5.16	5.97	5.97				
15	PLACE1001771	1.84	1.98	1.82	2.36	2.85	5.41	2.31	1.87	1.87				
	PLACE1001775	1.14	0.68	0.37	2.02	1.85	1.82	2.01	0.97	0.97	**	+		
	PLACE1001777	17.14	13.64	18.62	21.05	26.38	21.12	40.01	76.23	76.23		*	+	
	PLACE1001781	2.45	1.71	2.59	2.44	2.81	2.52	2.91	5.33	5.33				
	PLACE1001783	4.43	2.58	2.66	2.32	3.33	2.65	2.54	4.19	4.19				
20	PLACE1001786	1.74	1.05	1.30	1.23	1.66	1.40	1.26	1.69	1.69				
	PLACE1001788	5.13	2.94	2.51	5.8	4.90	5.17	4.40	3.27	3.27				
	PLACE1001795	2.72	1.91	2.58	4.69	4.12	5.43	5.56	6.85	6.85	**	+	**	+
	PLACE1001799	3.74	3.45	3.29	3.65	3.39	3.75	3.22	5.05	5.05				
	PLACE1001810	2.43	0.99	1.08	2.55	2.52	2.29	2.26	1.22	1.22				
25	PLACE1001817	6.6	4.05	4.21	9.77	8.48	6.29	8.47	8.36	8.36		*	+	
	PLACE1001821	3.26	2.45	2.55	4.22	4.44	5.51	4.69	7.27	7.27	*	+	*	+
	PLACE1001836	4.29	2.26	1.81	2.56	3.00	3.57	2.41	2.93	2.93				
	PLACE1001844	1.78	2.16	1.61	2.8	3.57	4.27	2.87	4.20	4.2	*	+	*	+
	PLACE1001845	2.41	1.41	2.18	4.39	5.00	4.06	2.82	2.33	2.33	**	+		
	PLACE1001858	4.51	4.42	4.15	7.53	6.22	8.84	4.27	3.55	3.55	*	+		
30	PLACE1001869	3.09	2.60	2.08	2.74	2.72	3.73	1.99	3.40	3.4				
	PLACE1001890	2.77	2.42	1.39	7.46	6.18	5.66	5.49	5.13	5.13	**	+	**	+
	PLACE1001897	2.18	2.26	1.85	6.69	5.35	5.34	8.97	9.82	9.82	**	+	**	+
	PLACE1001902	31.17	17.00	21.61	32.58	37.84	31.63	15.20	15.90	15.9				
	PLACE1001904	3.92	3.02	3.25	2.81	3.73	3.19	4.96	4.49	4.49		*	+	
35	PLACE1001907	5.11	3.84	3.69	6.62	6.43	7.96	4.32	5.12	5.12	*	+		
	PLACE1001910	1.87	3.06	2.35	3.3	3.81	3.68	14.39	26.30	26.3	*	+	**	+
	PLACE1001912	2.63	0.79	1.20	4.38	3.77	3.71	2.02	2.67	2.67	*	+		
	PLACE1001918	10.38	7.15	8.90	11.66	9.55	15.16	10.15	14.11	14.11				
	PLACE1001920	2.53	1.11	1.05	1.68	3.07	1.48	1.79	0.84	0.84				
40	PLACE1001928	8.17	4.57	3.74	7.72	5.90	6.65	3.44	4.51	4.51				
	PLACE1001930	2.19	1.43	2.13	1.81	3.19	3.67	2.17	2.30	2.3				
	PLACE1001949	2.08	1.14	1.41	2.07	1.98	1.77	1.69	2.05	2.05				
	PLACE1001959	1.52	1.78	2.06	2.37	1.77	2.84	1.64	2.36	2.36				
	PLACE1001969	4.16	2.19	2.62	4.17	4.18	4.94	2.88	2.78	2.78				
45	PLACE1001974	9.4	3.65	4.39	13.34	9.00	13.23	6.71	10.90	10.9				
	PLACE1001981	1.69	1.37	1.20	2.64	1.90	2.12	1.52	1.67	1.67	*	+		
	PLACE1001983	5.62	5.76	3.72	4.29	4.58	4.62	6.62	4.70	4.7				
	PLACE1001989	5.11	2.90	3.88	7.82	6.79	6.73	3.99	4.04	4.04	*	+		
	PLACE1002004	8.3	4.91	5.56	11.8	13.71	13.04	6.33	7.42	7.42	**	+		
	PLACE1002008	14.39	6.47	3.72	18.67	18.94	18.14	8.81	8.95	8.95	*	+		
50	PLACE1002015	8.41	6.18	4.44	7.71	7.18	8.85	8.17	7.96	7.96				
	PLACE1002044	1.09	1.29	1.71	3.03	2.76	2.81	3.27	3.06	3.06	**	+	**	+
	PLACE1002046	3.04	2.60	2.80	3.24	2.46	4.89	3.21	2.77	2.77				
	PLACE1002052	1.9	0.59	1.24	2.33	1.00	2.14	1.49	1.25	1.25				
	PLACE1002066	6.22	4.18	2.39	10.6	9.81	10.78	7.57	8.32	8.32	**	+	*	+
55	PLACE1002072	4.3	3.28	3.34	7.74	8.18	6.54	4.13	5.29	5.29	**	+		
	PLACE1002073	4.41	2.63	2.33	3.9	3.18	3.49	2.87	3.67	3.67				
	PLACE1002080	9.31	4.83	4.67	8.96	9.64	10.72	7.21	6.98	6.98				

Table 303

5	PLACE1001344	2.76	1.50	1.35	2.41	3.45	2.46	1.70	2.00	2				
	PLACE1001351	3.23	1.94	2.24	3.49	3.29	3.25	2.62	4.03	4.03				
	PLACE1001366	4.38	2.83	2.63	5.26	5.03	5.59	4.18	3.48	3.48	*	+		
	PLACE1001377	2.21	0.95	1.13	1.75	2.13	2.07	1.20	1.68	1.68				
	PLACE1001383	3.71	1.90	1.47	3.95	6.26	1.71	1.64	2.49	2.49				
10	PLACE1001384	3.18	2.05	1.78	4.94	5.31	4.83	2.21	2.83	2.83	**	+		
	PLACE1001387	4.38	2.11	2.54	3.04	2.86	4.24	2.34	3.05	3.05				
	PLACE1001395	1.59	1.26	1.15	3.65	3.08	5.18	3.82	2.99	2.99	*	+	**	+
	PLACE1001399	11.87	6.31	8.20	17.43	15.28	22.75	13.01	12.96	12.96	*	+		
	PLACE1001401	1.52	0.25	1.01	1.14	0.80	1.79	1.18	1.33	1.33				
	PLACE1001407	6.8	4.32	5.87	3.76	3.93	5.36	10.73	10.24	10.24		**	+	
15	PLACE1001412	5.12	1.76	2.22	3.71	2.25	2.65	2.13	1.31	1.31				
	PLACE1001414	15.81	9.44	8.70	18.1	13.15	13.80	12.97	12.27	12.27				
	PLACE1001416	4.85	3.13	3.24	4.86	3.47	4.68	3.85	4.04	4.04				
	PLACE1001433	34.75	27.32	25.94	41.44	46.72	44.79	20.21	24.82	24.82	**	+		
	PLACE1001440	3.36	1.52	3.50	3.58	3.41	4.36	3.30	2.97	2.97				
20	PLACE1001456	2.82	2.23	1.05	4.35	4.43	4.27	3.77	3.38	3.38	*	+		
	PLACE1001464	1.12	0.36	0.61	1.11	1.20	1.53	4.05	3.36	3.36		**	+	
	PLACE1001468	1	1.48	0.93	1.65	1.22	1.79	1.02	0.92	0.92				
	PLACE1001484	5.54	3.35	3.73	7.43	7.35	10.20	3.71	4.16	4.16	*	+		
	PLACE1001500	8.54	6.02	4.38	7.39	7.18	5.61	5.36	6.08	6.08				
25	PLACE1001502	6.06	4.35	3.12	4.46	5.05	4.69	4.11	4.84	4.84				
	PLACE1001503	6.09	4.19	3.41	7.11	7.79	6.61	4.97	5.70	5.7	*	+		
	PLACE1001505	20.88	12.93	14.68	15.96	17.98	17.32	9.92	14.48	14.48				
	PLACE1001513	6.48	3.77	5.22	5.72	3.68	4.54	4.27	6.65	6.65				
	PLACE1001516	10.93	7.17	9.57	12.22	8.39	12.84	8.43	11.33	11.33				
	PLACE1001517	5.77	3.37	4.96	7.37	4.67	6.00	5.80	4.89	4.89				
30	PLACE1001523	23.41	10.77	16.66	12.24	9.55	12.27	10.99	12.94	12.94				
	PLACE1001526	7.32	4.41	2.62	6.04	11.01	4.64	4.47	5.72	5.72				
	PLACE1001534	4	1.96	2.04	4.38	6.28	3.78	3.64	3.03	3.03				
	PLACE1001536	2.83	1.23	1.62	1.76	3.23	2.47	2.13	1.81	1.81				
	PLACE1001545	36.23	12.22	23.79	37.99	57.83	39.02	33.62	43.32	43.32				
35	PLACE1001551	6.66	3.51	3.07	3.77	5.41	4.65	3.22	3.12	3.12				
	PLACE1001564	1.35	0.83	1.14	1.76	1.17	1.28	1.94	2.02	2.02		**	+	
	PLACE1001570	0.93	0.34	0.64	2.16	2.60	4.80	1.89	2.31	2.31	*	+	**	+
	PLACE1001571	7.95	4.12	4.74	8.82	11.30	11.21	6.14	8.15	8.15	*	+		
	PLACE1001595	11.96	8.35	6.84	10.3	8.39	8.08	8.16	6.97	6.97				
40	PLACE1001602	10.71	5.17	5.52	7.59	10.40	7.10	3.81	6.12	6.12				
	PLACE1001603	2.7	2.04	2.99	5.01	5.83	4.53	3.42	3.10	3.1	**	+		
	PLACE1001608	2.44	2.10	2.41	3.4	4.03	5.05	2.95	3.88	3.88	*	+	*	+
	PLACE1001610	5.43	4.80	5.73	13.88	9.92	13.14	7.65	8.25	8.25	**	+	**	+
	PLACE1001611	3.56	2.47	3.24	3.84	3.75	5.73	3.92	3.82	3.82				
45	PLACE1001629	6.48	3.78	4.26	6.9	3.97	6.33	1.49	1.62	1.62	*	+		
	PLACE1001632	8.49	4.09	6.12	10.6	10.02	12.44	6.25	8.12	8.12	*	+		
	PLACE1001634	3.06	1.40	1.54	5.61	6.76	4.47	2.48	4.23	4.23	*	+		
	PLACE1001637	4.89	3.35	2.51	2.97	3.72	3.26	3.61	4.38	4.38				
	PLACE1001640	6.92	2.46	2.49	7.67	8.87	5.84	4.54	7.69	7.69				
	PLACE1001655	3.46	2.83	2.76	2.95	2.85	2.93	2.12	2.26	2.26	*	+		
50	PLACE1001672	3.35	1.68	2.29	4.35	2.93	3.76	4.49	2.60	2.6				
	PLACE1001676	1.74	0.78	2.18	1.12	1.17	2.29	1.50	2.79	2.79				
	PLACE1001683	8.62	6.71	9.02	11.78	10.73	12.96	10.63	12.86	12.86	*	+	*	+
	PLACE1001691	5.26	3.63	4.10	12.07	10.05	6.33	3.77	5.16	5.16	*	+		
	PLACE1001692	4.42	2.12	2.27	4.86	5.36	4.90	4.07	3.26	3.26				
55	PLACE1001705	8.07	4.26	3.08	6.53	6.17	7.84	4.62	7.50	7.5				
	PLACE1001716	3.8	1.68	2.70	3.78	4.85	3.53	4.66	5.71	5.71	*	+		
	PLACE1001720	1.91	1.29	2.34	3.39	3.22	2.45	2.24	3.40	3.4	*	+		

Table 302

5	PLACE1000977	2.41	2.94	1.04	2.67	2.65	2.73	2.52	2.72	2.72				
	PLACE1000979	9.34	4.89	6.74	13.62	13.31	16.23	7.57	8.33	8.33	**	+		
	PLACE1000986	4.3	2.25	2.59	5.14	4.48	5.42	4.23	5.38	5.38	*	+		
	PLACE1000987	7.13	4.86	5.70	7.21	6.57	6.09	7.59	7.62	7.62				
	PLACE1001000	4.76	2.74	3.26	8.41	15.56	9.19	5.75	6.47	6.47	*	+	*	+
10	PLACE1001007	7.63	3.72	2.80	5.05	4.48	4.63	4.14	4.58	4.58				
	PLACE1001010	2.3	1.89	2.06	3.44	3.64	3.65	1.96	2.59	2.59	**	+		
	PLACE1001015	2.92	1.68	1.34	3.1	2.61	2.85	2.90	4.52	4.52				
	PLACE1001016	7.21	2.36	3.51	5.03	5.51	6.32	4.81	4.26	4.26				
	PLACE1001022	3.86	2.81	2.95	4.41	2.88	3.07	2.80	2.90	2.9				
15	PLACE1001024	3.88	2.20	3.13	2.3	2.95	4.59	2.73	3.68	3.68				
	PLACE1001036	5.16	2.56	3.47	6.09	4.65	5.59	4.01	4.38	4.38				
	PLACE1001038	28.81	14.88	16.16	21.4	17.66	19.48	21.32	28.28	28.28				
	PLACE1001048	3.36	1.96	1.23	2.27	1.42	1.71	1.83	3.38	3.38				
	PLACE1001054	7.9	5.99	5.59	6.24	6.31	4.84	4.36	6.39	6.39				
	PLACE1001062	7.2	5.87	4.94	11.02	9.95	11.12	6.47	7.34	7.34	**	+		
20	PLACE1001063	1.41	1.69	1.15	2.65	3.68	3.53	1.70	3.59	3.59	**	+		
	PLACE1001076	2.26	0.97	1.04	1.44	1.83	1.65	2.02	2.26	2.26				
	PLACE1001081	12.46	8.57	9.92	15.12	11.87	13.36	10.20	12.65	12.65				
	PLACE1001088	2.63	1.81	1.14	3.01	3.83	4.04	1.79	3.12	3.12	*	+		
	PLACE1001092	6.88	3.43	3.30	7.95	6.98	7.48	8.10	6.69	6.69				
25	PLACE1001098	3.19	4.37	2.61	7.39	7.22	4.69	3.98	5.42	5.42	*	+		
	PLACE1001100	4.67	2.56	3.28	9.14	7.82	8.01	4.36	9.43	9.43	**	+		
	PLACE1001104	4.42	3.38	3.50	3.41	4.47	4.62	3.50	5.47	5.47				
	PLACE1001114	6.37	3.02	3.19	9.14	6.05	8.38	4.84	6.58	6.58				
	PLACE1001118	8.99	8.41	8.16	18.03	15.27	17.69	9.35	8.27	8.27	**	+		
30	PLACE1001123	3.67	2.98	3.43	6.53	5.15	5.14	7.08	8.09	8.09	*	+	**	+
	PLACE1001136	6.74	4.90	3.41	11.43	11.92	9.20	6.63	6.95	6.95	*	+		
	PLACE1001144	5.3	3.83	2.70	9.8	6.14	5.78	3.32	5.22	5.22				
	PLACE1001147	6.12	3.41	3.43	6.85	6.67	6.42	5.03	6.28	6.28				
	PLACE1001148	3.16	1.95	1.69	2.9	2.48	3.03	1.39	4.13	4.13				
35	PLACE1001159	1.33	1.09	1.58	2.28	2.10	1.76	1.96	4.06	4.06	*	+	*	+
	PLACE1001168	1.82	0.78	1.16	1.62	1.75	2.87	2.70	3.06	3.06		**	+	
	PLACE1001171	2.35	1.34	1.61	1.46	3.10	2.35	2.90	1.94	1.94				
	PLACE1001183	1.79	2.36	1.72	2.21	1.23	3.26	2.19	2.54	2.54				
	PLACE1001185	5.46	4.74	4.40	6.41	7.88	5.56	6.42	6.55	6.55		**	+	
	PLACE1001201	6.18	4.83	3.75	5.34	5.15	4.77	3.30	2.90	2.9				
40	PLACE1001229	9.82	5.35	4.18	9.24	5.15	7.25	8.28	8.97	8.97				
	PLACE1001231	9.55	4.73	5.18	5.83	6.30	4.83	3.56	5.51	5.51				
	PLACE1001238	5.01	3.11	3.77	6.38	5.60	5.68	4.47	4.58	4.58	*	+		
	PLACE1001241	2.02	1.58	1.43	2.15	2.00	2.71	2.20	2.62	2.62		*	+	
	PLACE1001242	20.17	17.27	18.47	18.81	15.90	19.20	22.68	25.15	25.15		**	+	
45	PLACE1001247	9.52	6.34	6.64	10.32	8.10	12.11	5.62	8.24	8.24				
	PLACE1001250	3.73	2.44	3.14	5.42	5.19	6.97	2.55	3.26	3.26	*	+		
	PLACE1001257	6.68	5.11	2.77	13.34	10.34	12.61	4.57	6.16	6.16	**	+		
	PLACE1001272	6.36	4.10	3.44	7.49	6.51	5.32	5.11	6.14	6.14				
	PLACE1001279	2.31	1.92	1.89	3.68	2.64	2.53	2.56	2.17	2.17				
	PLACE1001280	2.63	3.05	1.70	3.8	3.92	4.33	2.23	2.86	2.86	*	+		
50	PLACE1001294	1.16	0.01	1.04	3.47	4.82	2.72	5.00	5.65	5.65	*	+	**	+
	PLACE1001295	4.29	3.95	3.46	3.47	3.85	3.06	4.31	5.32	5.32				
	PLACE1001300	2.58	2.11	2.36	2.54	2.70	1.93	2.20	3.72	3.72				
	PLACE1001304	6.77	5.82	8.24	17.77	13.08	18.60	7.86	9.34	9.34	**	+		
	PLACE1001311	5.16	4.21	2.93	10.01	8.67	7.40	5.99	7.20	7.2	**	+	*	+
55	PLACE1001323	7.17	3.76	3.29	11.33	10.13	10.10	6.13	5.77	5.77	*	+		
	PLACE1001325	2.41	1.50	1.58	5.07	4.29	3.56	1.94	2.67	2.67	**	+		
	PLACE1001340	8.91	4.41	6.17	8	8.54	6.15	5.24	8.59	8.59				

Table 301

	PLACE1000546	3.29	1.94	2.26	2.05	2.11	2.19	2.74	2.01	2.01				
5	PLACE1000547	5.79	5.41	5.37	8.99	6.38	9.62	5.74	8.94	8.94	*	+		
	PLACE1000560	3.31	3.53	2.48	3.26	3.84	4.27	3.25	2.77	2.77				
	PLACE1000562	5.48	3.54	4.16	6.47	7.13	6.86	5.29	6.77	6.77	*	+		
	PLACE1000564	2.28	2.89	3.32	2.89	4.25	5.04	4.28	3.71	3.71		*	+	
	PLACE1000583	10.76	7.63	6.51	18.65	16.27	17.87	10.12	7.24	7.24	**	+		
10	PLACE1000587	7.2	4.11	4.88	9.4	11.04	9.29	6.85	6.39	6.39	*	+		
	PLACE1000588	7.89	4.98	4.13	9.54	8.74	6.18	7.91	6.38	6.38				
	PLACE1000596	7.64	7.46	10.08	8.78	8.56	8.98	4.59	7.82	7.82				
	PLACE1000599	5.52	4.56	3.15	8.04	7.54	8.14	4.12	5.23	5.23	**	+		
	PLACE1000605	4.13	3.66	3.53	4.62	5.26	5.10	5.59	5.89	5.89	**	+	**	+
15	PLACE1000610	3.95	3.19	2.63	4.04	4.12	4.83	3.09	3.87	3.87				
	PLACE1000611	1.33	4.36	3.21	2.64	5.18	3.62	3.25	4.05	4.05				
	PLACE1000626	3.93	3.49	2.73	5.31	3.91	4.11	4.05	3.66	3.66				
	PLACE1000633	2.72	3.21	2.28	6.49	6.56	3.99	3.45	2.66	2.66	*	+		
	PLACE1000636	2.12	1.92	1.69	2.35	1.07	2.86	1.27	1.58	1.58				
20	PLACE1000653	2.8	1.22	1.84	2.02	2.53	1.75	1.81	4.26	4.26				
	PLACE1000656	9.31	7.34	8.14	10.31	10.47	9.31	12.81	14.00	14		**	+	
	PLACE1000663	1.27	0.67	0.99	1.89	1.74	1.74	1.26	1.65	1.65	*	+		
	PLACE1000706	11.24	11.57	11.40	19.1	16.63	21.24	10.14	12.25	12.25	**	+		
	PLACE1000712	1.84	3.33	4.09	6.55	4.54	5.89	4.64	6.19	6.19	*	+	*	+
25	PLACE1000716	2.94	0.83	1.14	1.67	1.91	1.48	1.97	1.39	1.39				
	PLACE1000740	3.04	1.05	2.32	2.9	2.88	3.09	2.84	2.88	2.88				
	PLACE1000748	6.27	3.34	3.42	5.4	6.40	3.86	2.84	3.25	3.25				
	PLACE1000749	12.36	6.45	8.51	10.43	9.17	13.07	10.01	13.44	13.44				
	PLACE1000751	2.38	1.17	1.02	4.52	3.07	2.68	4.21	4.95	4.95		**	+	
	PLACE1000755	2.51	1.55	1.57	3.46	3.45	4.83	2.60	2.77	2.77	*	+		
30	PLACE1000769	2.21	1.01	1.04	2.25	2.24	3.89	2.18	2.07	2.07				
	PLACE1000778	5.1	3.19	2.79	4.88	3.83	3.91	3.55	2.38	2.38				
	PLACE1000785	8.86	6.54	5.09	10.87	11.53	8.38	4.96	7.33	7.33				
	PLACE1000786	4.27	4.46	2.71	4.67	3.49	4.67	3.76	4.74	4.74				
	PLACE1000793	6.19	3.54	4.79	9.71	9.92	9.47	5.31	5.48	5.48	**	+		
35	PLACE1000795	9.72	4.72	5.55	4.52	4.48	3.39	4.32	4.67	4.67				
	PLACE1000798	1.9	1.59	2.33	3.4	3.26	3.47	1.64	2.26	2.26	**	+		
	PLACE1000812	2.3	2.38	1.85	3.32	3.27	4.96	2.41	3.24	3.24	*	+		
	PLACE1000823	7.01	4.40	5.61	12.77	10.75	11.18	7.00	5.92	5.92	**	+		
	PLACE1000825	6.13	3.73	3.27	7.05	6.77	5.20	4.28	5.79	5.79				
40	PLACE1000838	5.14	3.45	2.78	6.34	7.02	4.42	12.05	18.19	18.19		**	+	
	PLACE1000841	3.14	5.34	2.01	3.49	3.92	2.49	3.35	1.76	1.76				
	PLACE1000843	4.46	2.15	3.63	4.5	6.77	4.11	1.87	4.89	4.89				
	PLACE1000849	10.82	6.77	8.57	8.51	10.69	9.82	7.58	11.02	11.02				
	PLACE1000856	2.83	1.51	2.02	3.37	2.62	2.73	2.59	1.96	1.96				
45	PLACE1000863	9.64	6.13	6.86	5.2	5.82	6.39	5.18	5.81	5.81				
	PLACE1000876	7.89	4.38	5.88	7.14	5.51	8.48	7.94	7.18	7.18				
	PLACE1000899	3.08	2.81	1.69	4.08	4.67	3.67	3.31	2.41	2.41	*	+		
	PLACE1000907	16.44	10.14	7.86	22.19	25.12	16.66	7.95	11.86	11.86				
	PLACE1000909	3.62	1.21	1.15	2.54	4.35	1.92	1.98	2.37	2.37				
	PLACE1000912	6.9	3.41	4.10	5.35	5.89	5.24	4.38	4.49	4.49				
50	PLACE1000914	3.46	1.48	2.11	2.59	3.24	2.71	3.41	2.78	2.78				
	PLACE1000918	0.79	0.41	0.85	0.84	1.40	1.52	0.67	1.33	1.33				
	PLACE1000927	3.51	2.64	4.51	6.98	7.67	10.76	8.88	8.80	8.8	*	+	**	+
	PLACE1000931	2.76	1.60	7.19	4.08	3.69	6.22	3.38	2.86	2.86				
	PLACE1000944	2.02	1.08	0.51	4.48	5.07	3.55	3.07	1.96	1.96	**	+		
55	PLACE1000948	3.27	0.90	1.90	2.66	2.46	1.89	1.91	1.97	1.97				
	PLACE1000958	2.75	1.53	1.51	2.98	2.99	3.11	3.17	4.29	4.29		*	+	
	PLACE1000972	6.67	4.02	6.08	7.27	8.73	6.46	4.61	7.73	7.73				

Table 300

	PLACE1000101	2.3	2.12	2.61	4.62	5.45	5.31	2.54	3.96	3.96	**	+		
5	PLACE1000121	3.32	1.82	3.36	3.18	3.46	3.22	4.10	2.97	2.97				
	PLACE1000133	22.32	10.62	12.41	24.57	19.93	22.03	9.44	17.41	17.41				
	PLACE1000142	3.77	2.94	3.97	3.72	2.78	3.50	4.86	3.02	3.02				
	PLACE1000146	12.04	5.71	7.52	11.96	8.63	12.12	6.11	6.64	6.64				
	PLACE1000163	10.38	6.77	6.39	8.08	8.26	4.88	8.20	6.01	6.01				
10	PLACE1000172	2.38	1.36	0.47	1.68	3.26	0.78	1.42	1.36	1.36				
	PLACE1000181	4.66	3.09	3.18	5.69	5.41	5.62	4.15	4.68	4.68	*	+		
	PLACE1000184	1.13	1.00	1.41	4.73	6.35	6.17	4.40	7.01	7.01	**	+	**	+
	PLACE1000185	5.78	3.85	4.83	5.4	6.28	6.49	6.72	6.56	6.56		*	+	
	PLACE1000198	3.55	2.09	2.55	2.87	3.21	4.22	3.14	3.19	3.19				
15	PLACE1000213	2.64	0.86	1.73	2.98	2.54	2.75	2.24	2.31	2.31				
	PLACE1000214	5.38	1.32	2.03	4.15	4.37	5.82	3.05	3.50	3.5				
	PLACE1000220	5.9	3.44	1.89	3.73	2.84	3.93	2.23	3.16	3.16				
	PLACE1000231	18.42	11.77	9.30	14.94	14.15	14.87	11.91	14.48	14.48				
	PLACE1000236	5.6	2.94	3.19	6.04	6.27	4.87	5.66	5.87	5.87				
20	PLACE1000245	7.5	5.11	6.34	10.03	9.79	11.42	4.16	7.99	7.99	**	+		
	PLACE1000246	5.62	3.38	4.68	6.48	8.30	6.53	8.63	9.43	9.43	*	+	**	+
	PLACE1000258	15.61	9.21	10.26	23.89	20.68	20.66	9.91	13.07	13.07	*	+		
	PLACE1000288	2.41	2.18	2.21	2.88	1.68	2.31	2.41	3.07	3.07				
	PLACE1000292	5.99	4.40	5.17	20.8	17.62	19.45	12.37	20.25	20.25	**	+	**	+
	PLACE1000302	1.46	1.42	1.22	6.15	8.89	5.78	5.17	5.07	5.07	**	+	**	+
25	PLACE1000304	4.47	1.71	1.91	3.89	2.76	3.12	2.80	2.80	2.8				
	PLACE1000308	4.91	2.41	1.59	3.39	5.24	3.59	2.01	2.78	2.78				
	PLACE1000309	11.75	7.68	5.52	7.14	11.13	6.51	7.34	11.09	11.09				
	PLACE1000312	4.15	1.12	1.95	3.37	3.51	3.75	2.70	2.85	2.85				
	PLACE1000330	2.07	1.35	1.92	2.05	1.50	2.72	2.22	2.82	2.82		*	+	
30	PLACE1000332	0.54	0.37	0.59	1.08	1.22	2.14	1.43	1.37	1.37	*	+	**	+
	PLACE1000347	3.56	1.98	2.82	5.26	6.11	4.66	4.59	5.16	5.16	*	+	*	+
	PLACE1000351	5.67	4.34	5.42	8.3	7.13	5.49	4.92	6.31	6.31				
	PLACE1000374	9.15	6.32	6.28	12.33	8.13	8.69	5.60	5.63	5.63				
	PLACE1000380	8.21	2.59	3.63	4.88	6.57	5.07	4.83	5.60	5.6				
35	PLACE1000383	3.43	2.31	1.31	2.37	3.17	2.14	2.59	1.96	1.96				
	PLACE1000397	4.72	2.15	2.60	3.29	2.51	3.41	2.52	3.33	3.33				
	PLACE1000401	8.18	4.62	4.15	5.55	6.29	6.94	5.61	6.88	6.88				
	PLACE1000406	5.56	3.08	2.60	5.54	5.34	5.46	3.82	3.45	3.45				
	PLACE1000412	3.31	2.01	1.64	4.18	4.67	3.93	2.55	2.54	2.54	*	+		
40	PLACE1000420	10.38	5.91	5.93	8.64	10.82	10.12	5.86	5.89	5.89				
	PLACE1000421	3.59	3.04	2.31	4.45	4.08	3.36	2.89	3.75	3.75				
	PLACE1000423	2.95	2.15	1.93	20.49	20.83	20.84	13.81	14.04	14.04	**	+	**	+
	PLACE1000424	3	2.12	1.66	4.43	3.32	3.59	1.60	2.52	2.52	*	+		
	PLACE1000430	3.63	1.51	1.58	2.45	2.43	3.11	1.57	3.03	3.03				
	PLACE1000433	4.59	1.89	2.39	2.55	2.63	3.39	3.84	2.91	2.91				
45	PLACE1000435	4.53	3.13	3.03	9.09	8.39	8.75	5.45	3.19	3.19	**	+		
	PLACE1000437	2.55	2.34	2.51	7.65	7.02	9.50	7.52	8.64	8.64	**	+	**	+
	PLACE1000442	12.33	5.94	10.64	23.09	26.07	18.20	10.78	10.42	10.42	*	+		
	PLACE1000444	9.31	6.03	5.50	16.99	19.05	17.38	8.02	10.01	10.01	**	+		
	PLACE1000453	6.66	4.79	5.00	7.58	6.74	7.37	6.15	9.05	9.05				
50	PLACE1000456	4.25	3.10	2.24	3.67	3.13	4.02	3.33	4.78	4.78				
	PLACE1000465	5.73	3.62	3.38	4.99	3.38	5.47	5.67	4.76	4.76				
	PLACE1000481	5.42	4.78	5.17	5.8	8.48	10.90	5.11	5.58	5.58				
	PLACE1000492	4.42	2.55	3.57	3.46	5.78	6.28	4.30	4.90	4.9				
	PLACE1000508	4.11	3.53	2.58	3.28	3.70	3.99	2.37	3.96	3.96				
55	PLACE1000512	5.22	2.40	1.36	6.14	5.78	4.97	4.87	4.74	4.74				
	PLACE1000540	2.6	2.41	1.99	4.78	4.15	4.34	1.97	1.70	1.7	**	+		
	PLACE1000541	6.4	6.38	5.54	8.78	7.96	6.93	7.44	11.12	11.12	*	+	*	+

Table 299

5	OVARC1001916	6.98	5.21	4.19	6.6	6.42	9.56	6.23	7.95	7.95				
	OVARC1001928	2.06	0.85	1.79	2.38	2.75	2.84	3.26	4.05	4.05			**	+
	OVARC1001937	3.08	3.56	3.08	6.71	6.67	8.66	8.49	10.57	10.57	**	+	**	+
	OVARC1001940	2.73	1.83	2.29	2.9	3.41	3.46	2.76	3.64	3.64	*	+		
	OVARC1001942	7.33	6.50	6.76	5.22	5.72	6.21	4.66	4.79	4.79	*	-	**	-
	OVARC1001943	10.42	8.83	6.98	5.68	5.59	6.08	6.06	4.31	4.31	*	-	*	-
10	OVARC1001949	10.36	7.25	8.90	17.76	16.95	14.34	7.64	7.78	7.78	**	+		
	OVARC1001950	6.51	3.98	3.61	6.85	7.18	5.80	4.70	6.17	6.17				
	OVARC1001952	8.93	7.35	6.04	9.34	7.56	8.32	8.80	9.41	9.41				
	OVARC1001954	2.25	1.93	2.80	2.22	2.67	3.53	3.43	3.10	3.1			*	+
	OVARC1001963	4.35	4.65	3.70	6.06	6.92	7.14	5.20	5.61	5.61	**	+	*	+
15	OVARC1001983	14.69	9.15	11.07	15.77	13.57	18.65	18.62	19.08	19.08			*	+
	OVARC1001987	4.18	3.62	3.23	5.27	5.35	7.29	5.42	5.22	5.22	*	+	**	+
	OVARC1001989	4.53	2.66	2.25	6.48	8.72	7.41	3.80	4.09	4.09	*	+		
	OVARC1001991	10.96	5.93	5.69	9.46	8.32	6.27	7.05	6.60	6.6				
	OVARC1002005	5.4	3.75	4.99	8.51	8.21	8.60	5.67	7.46	7.46	**	+		
	OVARC1002044	5.75	6.74	4.12	8.85	9.04	10.30	6.19	6.78	6.78	*	+		
20	OVARC1002046	11.4	8.29	10.75	14.32	15.39	13.03	16.29	16.11	16.11	*	+	**	+
	OVARC1002050	7.01	4.34	4.11	5.04	4.91	6.69	6.80	8.61	8.61				
	OVARC1002058	2.46	2.25	3.14	3.04	3.77	4.08	4.59	3.85	3.85			*	+
	OVARC1002066	3.19	1.93	3.61	3.32	2.98	4.14	5.23	6.90	6.9			*	+
	OVARC1002082	4.87	5.01	3.84	11.38	12.17	13.39	6.27	6.19	6.19	**	+	*	+
25	OVARC1002091	9.15	5.09	5.80	7.51	5.64	6.50	4.50	6.13	6.13				
	OVARC1002092	1.08	0.92	1.01	1.95	2.31	1.47	1.26	2.01	2.01	*	+	*	+
	OVARC1002093	10.46	8.34	8.22	9.65	10.46	8.69	6.29	9.67	9.67				
	OVARC1002094	3.39	2.34	2.33	2.97	3.73	2.67	2.42	4.62	4.62				
	OVARC1002107	4.25	3.34	3.27	6.5	6.62	9.76	3.44	3.77	3.77	*	+		
30	OVARC1002112	10.9	8.09	8.28	16.78	13.09	25.94	13.30	14.51	14.51			**	+
	OVARC1002126	5.65	6.82	6.95	13.64	10.71	12.11	9.13	8.48	8.48	**	+	**	+
	OVARC1002127	2.58	2.03	3.02	3.02	3.11	2.31	3.36	3.37	3.37			*	+
	OVARC1002138	2.48	2.26	1.89	3.19	3.39	3.93	1.72	2.13	2.13	**	+		
	OVARC1002143	1.69	1.30	0.60	1.38	1.56	1.86	1.19	0.95	0.95				
35	OVARC1002156	1.66	0.93	0.95	1.52	1.87	1.95	2.12	1.74	1.74				
	OVARC1002158	2.7	2.62	1.87	2.12	2.65	2.44	2.26	2.68	2.68				
	OVARC1002165	7.2	5.63	4.73	11.72	8.43	11.59	6.50	7.88	7.88	*	+		
	OVARC1002176	8	8.96	7.89	12.99	11.14	15.46	14.15	11.02	11.02	*	+	*	+
	OVARC1002178	1.22	1.02	1.19	6.91	5.74	6.72	4.31	4.39	4.39	**	+	**	+
40	OVARC1002182	2.89	1.94	1.74	3.43	2.78	3.06	2.40	2.34	2.34				
	OVARC1002185	3.07	1.87	2.74	2.77	3.03	2.27	3.08	3.27	3.27				
	PLACE1000004	4.13	1.50	2.40	4.62	3.84	3.14	1.43	2.34	2.34				
	PLACE1000005	1.35	0.94	1.81	2.1	2.21	3.64	1.75	1.86	1.86				
	PLACE1000006	3.24	3.13	3.46	5.32	4.20	5.06	3.54	4.19	4.19	*	+	*	+
45	PLACE1000007	3.52	1.48	1.95	2.76	2.50	3.15	1.95	2.86	2.86				
	PLACE1000014	4.25	3.03	3.71	8.86	8.24	8.01	5.81	6.21	6.21	**	+	**	+
	PLACE1000031	2.43	0.83	0.85	3.06	2.75	3.91	2.27	1.91	1.91	*	+		
	PLACE1000033	1.29	0.90	0.41	1.55	1.06	1.17	1.59	1.10	1.1				
	PLACE1000040	4.49	2.71	2.01	6.89	9.12	6.89	4.66	5.42	5.42	*	+		
	PLACE1000048	1.6	1.02	1.34	5.06	4.76	4.04	3.48	3.87	3.87	**	+	**	+
50	PLACE1000050	5.68	3.49	4.13	5.18	4.97	6.58	3.56	3.95	3.95				
	PLACE1000061	158.3	101.17	90.85	157.97	122.81	120.53	58.82	94.38	94.38				
	PLACE1000066	24.72	10.40	14.31	13.08	14.83	12.97	11.89	17.52	17.52				
	PLACE1000075	3.77	2.50	2.49	11.38	15.88	19.81	6.47	10.82	10.82	**	+	*	+
	PLACE1000078	3.4	1.72	2.20	4.82	4.89	6.42	2.94	3.88	3.88	*	+		
55	PLACE1000081	10.27	4.42	4.34	7.73	8.15	4.92	5.06	4.85	4.85				
	PLACE1000086	7.07	5.86	4.84	7.21	7.07	4.90	5.17	6.83	6.83				
	PLACE1000094	3.81	2.40	2.03	2.26	2.48	2.45	2.38	2.04	2.04				

Table 298

5	OVARC1001555	6.13	2.98	2.93	3.66	4.13	4.35	3.24	3.51	3.51				
	OVARC1001560	5.27	2.89	4.00	3.57	5.47	3.00	1.86	5.44	5.44				
	OVARC1001569	4.31	1.79	2.67	5.77	3.68	6.02	3.66	4.73	4.73				
	OVARC1001570	3.15	1.30	2.66	3.39	3.35	3.15	3.14	2.39	2.39				
	OVARC1001577	4.77	2.77	4.00	5.05	6.04	4.74	3.79	3.40	3.4				
	OVARC1001578	0.13	0.13	0.49	0.11	0.08	0.34	(0.16)	0.33	0.33				
10	OVARC1001596	6.65	4.15	4.07	12.92	13.04	11.27	13.75	17.88	17.88	**	+	**	+
	OVARC1001600	4.44	1.10	1.82	4.64	5.45	5.21	2.46	3.26	3.26				
	OVARC1001607	3.4	1.49	1.81	4.77	3.07	3.12	3.27	4.29	4.29				
	OVARC1001610	1.98	0.84	1.36	1.63	3.05	2.07	1.29	1.68	1.68				
	OVARC1001611	2.19	0.50	1.35	1.78	1.02	1.32	1.66	1.19	1.19				
15	OVARC1001615	4.22	1.84	2.90	5.28	3.15	3.01	2.44	2.96	2.96				
	OVARC1001636	1.51	1.25	1.84	2.49	2.09	2.98	2.73	3.68	3.68	*	+	**	+
	OVARC1001668	12.16	5.32	7.43	18.64	16.53	18.49	8.30	9.71	9.71	*	+		
	OVARC1001702	8.57	3.96	3.47	6.26	5.42	3.41	3.42	6.27	6.27				
	OVARC1001703	3.45	1.33	2.17	2.9	2.76	1.60	1.67	2.48	2.48				
20	OVARC1001710	12.16	6.40	8.14	12.51	12.10	10.06	5.91	10.48	10.48				
	OVARC1001711	3.85	1.19	3.00	4.46	4.77	3.21	3.17	3.47	3.47				
	OVARC1001713	3.83	1.81	3.06	4	3.01	2.37	3.41	2.97	2.97				
	OVARC1001725	1.76	0.84	1.52	1.59	1.72	1.08	1.90	2.27	2.27				
	OVARC1001726	5.39	1.55	3.13	5.82	3.63	5.08	3.26	3.16	3.16				
25	OVARC1001727	0.29	0.42	1.02	0.81	1.66	2.65	0.38	0.85	0.85				
	OVARC1001731	69.09	38.65	38.62	61.15	63.80	29.40	50.44	54.36	54.36				
	OVARC1001735	3.44	1.71	2.00	2.93	3.19	1.89	1.63	2.09	2.09				
	OVARC1001741	5.73	2.80	4.04	7.5	7.39	7.90	7.54	6.67	6.67	*	+	*	+
	OVARC1001745	7.24	4.36	4.49	8.97	10.22	8.41	6.60	5.98	5.98	*	+		
30	OVARC1001759	1.01	0.86	1.04	1.08	1.84	2.94	2.19	2.25	2.25		**	+	
	OVARC1001762	8.58	3.74	6.34	5.15	5.47	7.03	4.95	5.82	5.82				
	OVARC1001766	9.38	4.99	6.59	7.66	8.01	9.59	6.94	8.67	8.67				
	OVARC1001767	3.53	1.57	1.68	5.51	3.61	4.66	1.50	1.77	1.77	*	+		
	OVARC1001768	2.87	1.10	1.41	3.92	5.14	2.20	2.97	2.24	2.24				
	OVARC1001770	8.73	3.17	3.93	4.79	3.74	3.92	3.08	5.26	5.26				
35	OVARC1001776	9.28	3.35	3.86	7.43	6.75	3.40	4.83	5.46	5.46				
	OVARC1001791	6.37	2.23	2.37	4.77	4.93	3.53	3.51	5.12	5.12				
	OVARC1001795	3.33	1.66	2.08	3.57	2.56	3.39	2.70	4.38	4.38				
	OVARC1001798	7.18	6.07	6.66	13.95	10.63	12.79	7.22	8.63	8.63	**	+		
40	OVARC1001802	9.19	4.54	5.70	10.35	10.30	12.39	7.34	10.40	10.4	*	+		
	OVARC1001805	4.64	2.74	4.36	2.74	2.72	4.62	3.49	2.65	2.65				
	OVARC1001807	8.77	5.93	4.12	6.55	5.33	4.82	5.91	7.39	7.39				
	OVARC1001809	6.83	4.86	4.27	6.09	6.40	3.73	5.14	5.48	5.48				
	OVARC1001812	4.12	3.13	3.09	7.67	7.95	5.93	3.66	6.68	6.68	**	+		
	OVARC1001813	5.43	3.76	2.36	6.97	8.29	5.75	4.00	5.14	5.14				
	OVARC1001820	5.44	2.59	2.92	7.68	8.81	9.74	4.50	3.53	3.53	**	+		
45	OVARC1001828	1.52	0.56	0.82	0.49	1.38	1.06	0.77	2.57	2.57				
	OVARC1001833	6.47	2.16	4.12	4.91	4.44	4.92	4.40	5.06	5.06				
	OVARC1001839	3.71	1.97	2.01	2.39	2.11	1.77	2.84	1.57	1.57				
	OVARC1001846	4.41	2.73	3.00	4.53	4.51	2.44	2.43	1.95	1.95				
	OVARC1001849	7.54	4.93	4.04	7.29	7.04	10.00	6.63	6.98	6.98				
50	OVARC1001861	6.18	3.30	3.37	5.23	6.05	5.82	6.62	5.17	5.17				
	OVARC1001873	2.23	3.58	2.82	5.06	4.34	4.98	4.48	5.41	5.41	*	+	*	+
	OVARC1001879	6.45	3.48	3.55	6.19	6.28	6.46	4.62	5.20	5.2				
	OVARC1001880	8.1	5.60	6.83	9.11	8.57	12.18	8.27	7.92	7.92				
	OVARC1001883	2.85	1.41	1.74	4.9	4.51	4.19	2.29	2.05	2.05	**	+		
55	OVARC1001900	4.98	3.20	2.77	3.89	3.38	3.75	3.72	2.89	2.89				
	OVARC1001901	4.87	3.60	3.92	3.84	3.21	3.00	1.68	3.04	3.04				
	OVARC1001911	6	4.01	3.43	3.55	3.02	2.97	2.70	4.72	4.72				

Table 297

	OVARC1001243	1.72	1.35	1.37	1.54	2.52	1.73	1.36	2.41	2.41			
5	OVARC1001244	24.7	9.04	13.89	22.81	23.41	15.18	12.84	15.77	15.77			
	OVARC1001246	40.74	22.08	30.73	92.94	72.86	54.67	53.93	71.88	71.88	*	+	*
	OVARC1001247	8.36	4.54	5.70	8.31	7.58	6.86	6.44	6.70	6.7			
	OVARC1001260	5.56	1.98	3.43	3.72	4.11	5.56	3.81	5.29	5.29			
	OVARC1001261	7.49	5.34	5.88	8.27	8.14	6.50	4.18	3.66	3.66		*	-
10	OVARC1001268	9.66	6.34	6.78	20.35	19.09	14.70	18.61	12.90	12.9	**	+	*
	OVARC1001270	2.46	0.92	1.16	1.01	0.99	1.69	1.24	1.98	1.98			
	OVARC1001271	7.39	3.05	5.29	8.27	10.72	9.05	7.37	6.66	6.66	*	+	
	OVARC1001282	1.01	0.92	0.97	0.97	2.26	1.76	1.02	2.02	2.02			
	OVARC1001296	2.46	1.56	1.43	2.56	2.90	3.81	2.32	2.50	2.5			
15	OVARC1001306	7.3	3.30	5.02	6.03	4.37	5.50	5.45	6.39	6.39			
	OVARC1001314	0.91	0.46	0.79	1.37	1.95	2.32	1.59	1.62	1.62	*	+	**
	OVARC1001316	1.39	0.64	0.79	0.83	1.74	1.83	1.60	1.04	1.04			
	OVARC1001329	14.48	8.75	10.68	26.47	22.48	16.87	10.91	14.31	14.31	*	+	
	OVARC1001330	5.69	3.01	1.92	3.71	3.31	3.24	2.35	2.85	2.85			
	OVARC1001336	5.35	4.02	3.78	4.8	5.04	6.17	4.16	5.22	5.22			
20	OVARC1001338	3	2.42	3.08	2.63	3.26	3.21	2.60	4.03	4.03			
	OVARC1001339	18.39	11.67	11.13	15.76	12.03	15.86	13.83	17.02	17.02			
	OVARC1001340	3.7	2.44	2.40	2.48	2.50	2.72	1.64	1.40	1.4		*	-
	OVARC1001341	9.61	7.33	5.62	10.7	12.45	13.37	7.41	10.65	10.65	*	+	
	OVARC1001342	133.57	112.33	102.75	148.81	134.63	172.83	71.00	44.68	44.68		**	-
25	OVARC1001344	7.19	4.91	4.20	12.04	11.73	10.02	5.70	6.29	6.29	**	+	
	OVARC1001357	1.77	0.51	0.85	0.71	1.22	1.30	1.05	2.71	2.71			
	OVARC1001359	12.91	9.14	12.19	10.45	11.07	11.24	11.72	11.75	11.75			
	OVARC1001360	1.13	0.79	1.43	0.68	1.47	0.77	1.27	2.96	2.96			
	OVARC1001369	3.18	3.27	2.79	3.55	2.73	3.58	3.69	3.39	3.39			
30	OVARC1001372	2.77	2.30	1.69	2.23	2.48	3.94	3.04	2.69	2.69			
	OVARC1001376	2.87	2.00	1.97	5.27	5.80	7.45	3.84	3.47	3.47	**	+	*
	OVARC1001381	9.02	7.72	5.78	16.38	17.31	19.84	9.24	7.41	7.41	**	+	
	OVARC1001391	4.51	2.73	2.85	3.51	4.11	3.13	3.49	3.91	3.91			
	OVARC1001392	8.74	6.58	5.89	10.76	13.40	11.71	12.35	14.18	14.18	*	+	**
35	OVARC1001399	8.85	5.58	4.72	7.92	8.25	8.82	4.81	5.40	5.4			
	OVARC1001417	2.7	1.43	2.23	1.21	1.52	2.52	2.51	2.99	2.99			
	OVARC1001419	4.3	5.24	4.00	3.68	3.86	6.94	5.84	6.00	6		*	+
	OVARC1001425	2.29	2.40	2.49	3.29	2.74	4.54	3.29	3.09	3.09		**	+
	OVARC1001436	2.31	2.50	1.77	3.81	3.30	4.11	3.38	2.41	2.41	**	+	
	OVARC1001442	3.28	3.48	2.35	2.21	3.99	4.48	3.98	3.31	3.31			
40	OVARC1001451	2.33	1.90	1.35	3.6	3.77	3.76	1.55	1.55	1.55	**	+	
	OVARC1001452	3.08	2.65	1.79	3.37	3.43	2.89	2.90	3.86	3.86			
	OVARC1001453	1.36	0.57	0.90	1.69	3.97	2.45	2.96	1.73	1.73			
	OVARC1001476	9.08	6.86	7.98	15.11	12.70	14.85	28.29	23.49	23.49	**	+	**
	OVARC1001480	2.63	2.84	2.87	3.18	2.98	4.97	4.13	4.00	4		**	+
45	OVARC1001489	0.44	0.69	0.81	2.69	2.29	3.27	1.10	4.03	4.03	**	+	
	OVARC1001493	1.25	1.74	1.87	2.29	2.11	2.40	3.16	2.54	2.54	*	+	*
	OVARC1001496	8.58	6.56	5.62	10.89	7.25	13.93	7.36	6.38	6.38			
	OVARC1001499	2.77	1.81	1.79	9.3	11.43	8.77	4.71	4.12	4.12	**	+	**
	OVARC1001506	6.8	3.72	2.93	7.69	8.55	5.52	2.68	3.49	3.49			
50	OVARC1001509	1.55	1.98	1.98	5.58	4.61	5.41	3.85	2.97	2.97	**	+	*
	OVARC1001510	1.71	1.36	1.70	2.6	2.03	1.38	0.87	1.95	1.95			
	OVARC1001516	4.33	2.50	2.28	4.35	4.41	5.42	3.48	4.66	4.66			
	OVARC1001525	1.15	0.47	0.25	2.5	1.57	2.82	0.67	0.87	0.87	*	+	
	OVARC1001542	5.12	4.17	4.21	9.27	7.88	7.75	6.26	7.55	7.55	**	+	**
55	OVARC1001544	5.06	4.31	3.88	10.18	11.66	10.16	4.91	5.47	5.47	**	+	
	OVARC1001546	4.58	2.37	2.41	4.24	4.04	3.16	5.16	4.17	4.17			
	OVARC1001547	3.14	1.52	1.67	3.77	5.22	4.03	1.93	2.91	2.91	*	+	

Table 296

5	OVARC1001000	10.01	7.69	7.61	19.45	23.56	18.96	10.07	11.74	11.74	**	+	*	+
	OVARC1001004	1.03	0.80	0.91	1.57	2.14	1.61	1.90	1.48	1.48	*	+	*	+
	OVARC1001010	1.8	1.08	0.56	1.62	1.36	2.03	1.35	1.40	1.4				
	OVARC1001011	3.43	2.88	3.13	3.51	3.30	4.55	2.89	3.10	3.1				
	OVARC1001030	38.32	24.93	30.71	46.79	41.55	50.96	53.76	59.72	59.72	*	+	**	+
	OVARC1001032	1.55	1.32	1.67	3.18	2.58	2.77	2.83	1.37	1.37	**	+		
10	OVARC1001034	2.4	1.70	2.13	3.14	3.10	3.44	2.01	2.66	2.66	**	+		
	OVARC1001038	12.68	9.34	7.92	11.12	13.30	12.41	6.75	6.49	6.49				
	OVARC1001040	8.91	6.59	4.66	14.02	14.04	19.13	7.93	7.81	7.81	*	+		
	OVARC1001041	6.31	3.56	4.31	8.16	10.01	10.61	5.62	4.95	4.95	*	+		
	OVARC1001044	1.81	1.80	2.22	2.71	2.48	2.79	2.22	2.94	2.94	*	+		
15	OVARC1001049	9.39	8.47	8.39	15.62	16.10	16.18	9.93	8.69	8.69	**	+		
	OVARC1001051	57.5	54.01	57.15	51.44	56.52	72.78	36.05	33.73	33.73			**	-
	OVARC1001054	1.32	1.27	1.50	2.46	1.80	2.94	1.81	1.58	1.58	*	+	*	+
	OVARC1001055	3.77	1.65	2.45	4.24	4.50	2.62	2.94	3.56	3.56				
	OVARC1001062	11.74	5.75	4.85	11.81	10.68	10.78	3.12	5.25	5.25				
20	OVARC1001065	1.99	1.18	1.96	2.64	2.00	1.58	1.32	1.86	1.86				
	OVARC1001068	6.51	2.07	3.30	4.91	4.25	4.95	3.64	6.26	6.26				
	OVARC1001072	9.32	6.54	7.65	10.21	8.94	9.18	6.17	9.88	9.88				
	OVARC1001073	3.46	0.94	2.36	3.97	3.24	3.42	2.17	2.06	2.06				
	OVARC1001074	1.75	0.40	1.35	1.71	2.05	2.60	0.86	1.25	1.25				
25	OVARC1001078	7.1	3.90	5.62	11.77	8.65	7.84	4.87	6.07	6.07				
	OVARC1001085	5.2	2.42	3.41	5.59	4.12	3.31	4.28	6.32	6.32				
	OVARC1001086	5.76	2.45	2.47	3.85	5.26	3.78	2.15	3.47	3.47				
	OVARC1001091	3.91	3.54	2.95	5.93	5.39	4.15	4.20	3.41	3.41	*	+		
	OVARC1001092	4.33	2.96	3.51	6.04	6.34	5.50	3.69	5.56	5.56	**	+		
	OVARC1001104	1.53	0.53	0.40	1.32	1.57	1.20	0.63	1.14	1.14				
30	OVARC1001107	9.82	5.46	6.15	6.8	4.45	8.10	6.28	6.79	6.79				
	OVARC1001113	4.68	3.14	2.92	4.82	4.00	4.79	2.64	3.74	3.74				
	OVARC1001117	6.69	2.96	3.38	8.53	8.56	12.29	4.84	6.35	6.35	*	+		
	OVARC1001118	8.12	5.06	4.70	11.61	11.15	10.02	5.36	7.35	7.35	*	+		
	OVARC1001125	18.96	12.37	9.61	15.08	18.61	12.67	4.50	5.26	5.26			*	-
35	OVARC1001129	5.21	3.98	5.45	6.68	4.55	3.29	2.17	3.47	3.47			*	-
	OVARC1001132	6.52	3.70	5.55	7.12	8.81	9.06	2.18	2.72	2.72	*	+	*	-
	OVARC1001138	16.11	12.56	10.50	16.95	13.15	17.48	16.11	18.55	18.55				
	OVARC1001141	5.54	2.36	3.55	4.59	3.46	4.09	3.37	5.02	5.02				
	OVARC1001154	5.08	2.38	3.52	7.23	5.71	6.41	6.14	7.71	7.71	*	+	*	+
40	OVARC1001161	5.7	2.64	4.14	8.62	7.37	7.00	3.80	4.51	4.51	*	+		
	OVARC1001162	7.21	3.90	4.19	8.88	8.61	6.05	4.92	4.79	4.79				
	OVARC1001163	8.43	4.40	4.84	6.45	6.12	5.05	5.16	8.27	8.27				
	OVARC1001167	6.39	2.75	3.96	9.57	10.93	6.52	6.33	5.84	5.84				
	OVARC1001169	2.12	0.82	1.00	1.91	2.68	3.48	1.25	1.14	1.14				
	OVARC1001170	5.03	2.13	3.01	9.37	9.52	8.69	6.25	6.17	6.17	**	+	*	+
45	OVARC1001171	13.87	7.94	9.22	17.9	10.00	17.22	8.13	7.41	7.41				
	OVARC1001173	6.07	4.57	5.00	13.94	11.16	14.09	6.22	8.18	8.18	**	+	*	+
	OVARC1001176	120.6	80.54	85.77	70.13	72.81	62.53	40.27	47.98	47.98			*	-
	OVARC1001180	11.62	7.41	6.61	16.44	18.19	11.30	11.76	9.48	9.48				
	OVARC1001188	6.48	2.62	2.03	5.02	7.63	5.54	3.93	4.15	4.15				
50	OVARC1001200	2.22	1.30	1.20	5.74	5.77	3.98	2.73	3.72	3.72	**	+	*	+
	OVARC1001202	7.54	4.54	7.94	10.59	9.48	9.04	5.59	7.44	7.44				
	OVARC1001206	4.56	1.77	2.35	5.27	2.32	4.40	4.18	3.01	3.01				
	OVARC1001209	5.41	4.08	4.25	4.84	4.05	4.40	5.08	5.10	5.1				
	OVARC1001219	2.78	1.08	2.61	2.53	1.72	2.02	0.89	2.02	2.02				
55	OVARC1001222	2.69	0.99	2.05	4.33	4.36	2.81	4.34	4.56	4.56			**	+
	OVARC1001232	6.79	3.23	4.22	10.1	7.69	5.83	3.19	6.06	6.06				
	OVARC1001240	5.42	2.74	3.04	7.44	8.60	6.15	4.85	5.14	5.14	*	+		

Table 295

5	OVARC1000741	7.47	3.93	4.05	6.71	8.34	4.11	5.58	6.82	6.82			
	OVARC1000746	2.7	1.49	1.95	3.42	4.80	3.86	2.21	3.03	3.03	*	+	
	OVARC1000764	9.15	7.18	6.73	6.1	5.81	7.27	6.23	6.93	6.93			
	OVARC1000769	1.96	2.22	1.65	4.18	3.56	4.40	2.93	2.93	2.93	**	+	**+
	OVARC1000771	3.36	1.52	2.49	4.38	3.35	3.58	3.00	4.34	4.34			
	OVARC1000773	223.93	75.55	197.24	131.33	115.24	132.74	69.02	82.73	82.73			
10	OVARC1000775	5.89	2.38	2.57	10.9	11.89	6.67	5.95	7.36	7.36	*	+	
	OVARC1000778	5.16	2.89	2.70	7.19	7.19	4.94	4.21	3.79	3.79			
	OVARC1000779	1.34	0.25	1.68	0.81	2.17	1.66	0.98	2.78	2.78			
	OVARC1000781	3.01	1.11	1.81	3.21	4.12	3.42	2.43	3.96	3.96			
	OVARC1000787	5.12	1.26	2.40	6.21	4.91	6.16	2.68	3.80	3.8			
15	OVARC1000789	17.92	12.51	11.26	12.68	11.30	14.18	7.52	8.71	8.71			
	OVARC1000800	10.27	6.21	6.25	13.32	11.12	11.87	8.07	9.42	9.42	*	+	
	OVARC1000802	3.94	1.53	1.34	4.85	5.51	3.97	3.28	3.23	3.23			
	OVARC1000810	7.31	2.74	2.89	9.23	8.19	6.66	4.42	6.46	6.46			
	OVARC1000811	4.94	1.49	1.98	3.69	5.14	3.20	2.80	3.11	3.11			
20	OVARC1000814	8.98	4.85	4.30	12.34	14.84	13.49	5.29	9.28	9.28	*	+	
	OVARC1000816	5.55	2.23	3.34	6.25	6.38	4.13	4.96	10.86	10.86			
	OVARC1000817	0.67	0.84	0.17	1.03	1.43	0.88	1.03	1.18	1.18			
	OVARC1000834	7.9	3.52	4.48	7.01	4.99	6.90	5.30	8.11	8.11			
	OVARC1000846	8.76	5.89	5.62	13.13	13.07	12.45	7.92	8.86	8.86	**	+	
25	OVARC1000850	4.55	4.35	3.79	5.06	4.86	6.51	5.09	5.93	5.93		*	+
	OVARC1000853	10.26	6.75	7.96	17.45	22.42	13.77	15.27	17.34	17.34	*	+	**+
	OVARC1000862	2.31	1.51	1.67	2.98	3.34	3.48	2.84	3.92	3.92	**	+	*
	OVARC1000873	5.08	3.94	3.56	7.67	7.81	9.71	8.49	9.22	9.22	**	+	**+
	OVARC1000875	13.15	7.32	6.94	10.33	8.49	11.65	7.63	12.92	12.92			
	OVARC1000876	3.56	1.95	2.71	3.83	2.75	3.80	2.91	3.90	3.9			
30	OVARC1000883	11.24	5.79	7.03	7.42	6.63	8.18	6.12	10.30	10.3			
	OVARC1000885	1.99	1.85	0.96	2.91	2.72	4.05	1.81	1.84	1.84	*	+	
	OVARC1000886	3.79	3.90	3.30	5.23	4.59	3.88	4.18	4.19	4.19		*	+
	OVARC1000890	16.12	9.23	8.22	13.23	13.98	12.06	8.12	8.78	8.78			
	OVARC1000891	9.14	4.58	8.52	6.77	7.63	5.67	3.14	5.82	5.82			
35	OVARC1000897	1.42	0.51	0.89	0.57	0.73	1.37	0.82	2.14	2.14			
	OVARC1000912	3.17	1.30	1.93	1.64	2.12	2.69	2.76	3.24	3.24			
	OVARC1000914	1.78	1.84	1.59	1.55	2.20	2.36	1.62	3.25	3.25			
	OVARC1000915	6.15	3.81	2.82	7.18	8.08	11.76	6.61	6.54	6.54			
	OVARC1000916	3.99	3.47	3.85	4.78	5.13	5.34	5.36	5.24	5.24	**	+	**+
40	OVARC1000924	3.43	2.14	2.20	3.95	5.25	6.94	3.56	2.87	2.87	*	+	
	OVARC1000928	2.3	1.45	1.90	2.94	5.21	2.82	3.59	4.30	4.3			**+
	OVARC1000936	2.24	1.71	1.41	6.64	6.60	5.33	2.20	4.51	4.51	**	+	
	OVARC1000937	4.37	3.36	3.65	4.37	4.24	5.18	4.25	6.07	6.07			
	OVARC1000945	6.45	6.49	5.55	5.48	7.78	7.98	6.71	7.64	7.64			
45	OVARC1000948	0.7	1.14	1.80	0.8	1.44	1.73	1.79	1.83	1.83			
	OVARC1000956	4.51	3.81	4.84	5.21	4.57	6.49	5.02	4.22	4.22			
	OVARC1000959	3.91	3.29	2.51	6.03	6.37	10.71	4.34	3.63	3.63	*	+	
	OVARC1000960	15.58	12.49	8.74	25.21	26.65	32.56	13.09	13.48	13.48	**	+	
	OVARC1000964	7.4	4.78	4.49	6.36	5.85	5.60	6.26	4.60	4.6			
	OVARC1000971	2.15	0.88	1.23	2.29	3.01	2.68	1.01	1.16	1.16	*	+	
50	OVARC1000975	3.59	2.39	2.50	3.12	3.77	3.56	3.45	3.78	3.78			
	OVARC1000976	1.43	1.10	0.67	1.47	1.70	1.61	1.96	1.76	1.76		*	+
	OVARC1000981	4.17	3.59	2.08	3.86	4.73	5.67	6.53	5.87	5.87		*	+
	OVARC1000982	5.28	3.08	4.53	3.99	3.64	7.26	3.94	5.23	5.23			
	OVARC1000984	2.89	2.32	3.02	4.83	4.31	6.06	3.98	4.72	4.72	*	+	**+
	OVARC1000995	6.28	3.58	3.62	10.03	11.01	12.54	6.34	8.81	8.81	**	+	*
55	OVARC1000996	2.44	0.93	1.29	3.23	3.01	2.60	1.59	2.15	2.15	*	+	
	OVARC1000999	13.81	7.04	7.16	18.71	16.83	17.82	9.89	10.20	10.2	*	+	

Table 294

	OVARC1000414	2.94	2.41	3.01	5.83	4.82	5.60	3.16	3.78	3.78	**	+	*	+
5	OVARC1000420	11.4	6.17	7.59	9.95	9.38	10.06	10.09	13.16	13.16				
	OVARC1000421	8.6	6.78	5.53	8.33	7.86	10.75	8.17	6.59	6.59				
	OVARC1000427	3.68	2.71	4.36	3.26	4.27	4.49	3.23	3.96	3.96				
	OVARC1000431	28.24	22.85	26.14	17.5	18.78	21.85	14.12	15.50	15.5	*	-	**	-
	OVARC1000437	4.74	2.97	4.16	5.12	6.15	7.20	4.22	6.60	6.6				
10	OVARC1000439	7.31	6.90	5.38	7.44	6.69	8.00	6.48	4.83	4.83				
	OVARC1000440	10.79	6.84	6.93	7.88	7.24	7.80	6.48	7.22	7.22				
	OVARC1000442	5.47	3.48	2.90	10.37	8.05	7.61	4.21	5.71	5.71	*	+		
	OVARC1000443	2.37	1.87	2.77	3.52	3.55	4.55	2.82	6.19	6.19	*	+		
	OVARC1000461	3.39	2.34	2.79	3.41	2.83	2.56	4.13	3.34	3.34				
15	OVARC1000465	4.49	3.75	4.70	4.65	4.57	4.49	3.93	2.86	2.86				
	OVARC1000466	5.63	3.82	4.46	5.01	4.97	7.62	6.00	5.12	5.12				
	OVARC1000467	3.64	2.33	2.91	3.88	3.66	4.53	4.40	4.32	4.32		*	+	
	OVARC1000470	4.4	2.42	1.89	7.76	7.31	7.37	4.36	3.86	3.86	**	+		
	OVARC1000473	5.77	6.12	12.59	5.13	4.08	6.65	4.72	6.17	6.17				
20	OVARC1000479	10.65	6.40	6.55	8.36	8.23	12.25	7.74	6.99	6.99				
	OVARC1000484	7.73	3.54	4.68	14.41	17.12	13.60	11.04	9.93	9.93	**	+	*	+
	OVARC1000486	3.13	1.48	1.74	5.56	5.39	7.63	4.10	3.04	3.04	**	+		
	OVARC1000496	0.32	0.95	1.13	0.23	0.59	1.74	1.38	0.85	0.85				
	OVARC1000528	0.79	1.22	1.43	1.76	1.97	2.08	2.17	1.68	1.68	*	+	*	+
	OVARC1000522	4.89	4.05	3.21	7.99	8.62	12.13	8.58	8.73	8.73	*	+	**	+
25	OVARC1000526	5.23	3.76	3.40	9.44	8.41	9.79	6.60	6.83	6.83	**	+	**	+
	OVARC1000529	8.29	5.03	3.79	8.43	8.08	7.91	6.33	6.00	6				
	OVARC1000533	13.85	10.76	9.50	10.46	10.65	9.69	10.80	10.74	10.74				
	OVARC1000543	2.14	1.23	0.78	1.99	1.06	1.67	1.34	1.95	1.95				
	OVARC1000550	3.95	2.99	2.96	3.41	5.27	5.08	3.89	3.69	3.69				
30	OVARC1000553	7.96	6.39	6.63	10.34	11.92	12.52	8.20	8.94	8.94	**	+	*	+
	OVARC1000556	2.91	2.73	2.33	4.64	4.36	4.57	3.30	5.76	5.76	**	+		
	OVARC1000557	1.8	2.00	2.08	3.66	2.89	3.58	2.75	2.23	2.23	**	+		
	OVARC1000561	5.49	5.12	4.27	12.79	11.21	12.35	4.34	6.50	6.5	**	+		
	OVARC1000564	11	4.97	4.49	6.39	9.03	5.47	5.12	5.72	5.72				
35	OVARC1000573	3.43	1.54	1.73	4.84	5.71	5.20	3.22	2.70	2.7	**	+		
	OVARC1000576	22.35	9.42	12.58	14.84	14.82	13.96	18.96	21.39	21.39				
	OVARC1000578	3.78	1.92	1.91	7.25	4.00	7.95	3.26	3.45	3.45	*	+		
	OVARC1000581	2.32	0.98	1.31	2.39	2.02	2.50	0.87	2.36	2.36				
	OVARC1000586	4.15	3.94	3.82	5.69	4.46	5.03	7.98	9.37	9.37	*	+	**	+
	OVARC1000588	3.09	2.32	2.34	6.24	5.07	6.64	3.10	4.00	4	**	+	*	+
40	OVARC1000605	3.48	1.27	1.57	3.94	3.34	1.96	2.17	3.54	3.54				
	OVARC1000622	16.94	7.82	7.29	28.21	27.34	23.72	13.10	15.48	15.48	*	+		
	OVARC1000636	7.07	3.14	2.94	8.06	7.46	5.78	4.15	5.40	5.4				
	OVARC1000640	1.93	1.10	2.17	2.95	3.95	2.11	1.86	2.87	2.87				
	OVARC1000649	6.55	3.47	4.45	4.81	4.28	3.96	4.42	5.10	5.1				
45	OVARC1000661	8.83	4.09	5.47	7.42	7.41	7.08	6.76	7.35	7.35				
	OVARC1000677	5.49	3.25	4.84	5.23	4.19	5.75	3.47	4.95	4.95				
	OVARC1000678	3.24	2.45	2.41	6.56	3.68	3.55	3.07	3.23	3.23				
	OVARC1000679	2.29	2.05	2.51	5.5	7.69	3.61	2.67	3.02	3.02	*	+	*	+
	OVARC1000681	3.04	1.58	2.55	2.83	3.65	2.16	1.27	3.32	3.32				
50	OVARC1000682	5.34	2.89	3.15	10.89	12.80	6.98	6.81	6.70	6.7	*	+	*	+
	OVARC1000689	6.35	2.64	5.24	6.82	5.05	3.15	3.89	5.86	5.86				
	OVARC1000700	4.87	2.36	3.84	6.43	6.33	5.87	5.58	3.71	3.71	*	+		
	OVARC1000703	6.09	5.10	4.50	10.85	8.68	9.16	5.32	6.24	6.24	**	+		
	OVARC1000722	6.99	3.21	3.22	7.7	5.28	6.60	3.85	5.25	5.25				
55	OVARC1000726	12.55	5.82	7.48	9.62	7.07	8.57	9.99	8.90	8.9				
	OVARC1000727	8.32	3.91	3.99	6.93	6.40	4.72	3.99	5.01	5.01				
	OVARC1000730	6.1	3.39	3.84	6.3	8.93	6.59	2.98	3.46	3.46				

Table 293

	OVARC1000109	10.73	4.48	6.00	9.44	8.48	8.37	6.70	8.07	8.07			
5	OVARC1000113	4.43	3.28	2.32	5.28	7.68	6.28	3.04	3.01	3.01	*	+	
	OVARC1000114	4.61	1.82	2.98	6.68	7.59	8.77	4.82	5.56	5.56	*	+	
	OVARC1000133	2.28	0.62	2.11	1.97	3.23	1.32	1.31	3.42	3.42			
	OVARC1000137	7.57	3.31	3.78	7.45	5.45	6.40	5.03	9.51	9.51			
	OVARC1000139	8.5	5.04	5.90	7.42	5.19	7.20	5.43	7.04	7.04			
10	OVARC1000145	1.66	0.51	1.26	2.03	2.15	2.60	1.95	1.96	1.96	*	+	
	OVARC1000148	13.99	5.79	5.64	16.54	19.40	9.14	7.33	8.83	8.83			
	OVARC1000151	5.62	2.25	3.47	4.79	5.94	4.15	4.17	6.14	6.14			
	OVARC1000157	5.78	3.92	3.63	20.18	23.53	19.12	7.05	10.69	10.69	**	+	*
	OVARC1000162	1.04	0.27	1.30	1.82	2.05	0.82	1.71	1.67	1.67			
15	OVARC1000168	6.93	3.43	5.38	9.14	7.70	8.50	5.44	8.50	8.5	*	+	
	OVARC1000169	20.78	9.01	10.52	18.85	14.31	18.81	15.67	26.42	26.42			
	OVARC1000178	6.27	4.19	5.21	6.05	5.93	6.06	4.30	5.93	5.93			
	OVARC1000182	1.08	0.33	0.60	3.18	1.53	2.07	1.58	1.16	1.16	*	+	
	OVARC1000186	11.87	6.09	4.34	4.72	8.03	4.57	4.49	8.00	8			
20	OVARC1000188	6.88	3.30	4.11	6.26	4.11	4.48	4.18	5.80	5.8			
	OVARC1000191	2.39	0.93	1.25	1.87	4.24	1.53	1.02	3.43	3.43			
	OVARC1000198	7.48	2.50	4.22	12.55	13.51	9.27	4.79	6.14	6.14	*	+	
	OVARC1000208	7.66	5.85	6.85	11.11	11.76	10.78	8.71	6.63	6.63	**	+	
	OVARC1000209	5.19	2.21	3.10	4.98	5.19	3.99	3.67	6.12	6.12			
25	OVARC1000212	7.76	3.64	5.91	6.62	4.86	7.78	4.09	6.97	6.97			
	OVARC1000216	1.71	1.54	1.80	2.95	1.87	2.06	1.88	2.20	2.2		*	+
	OVARC1000240	9.19	4.82	3.93	10.89	11.55	7.32	4.66	6.08	6.08			
	OVARC1000241	8.4	2.88	3.50	6.97	5.95	3.69	4.83	5.66	5.66			
	OVARC1000249	5.89	2.71	3.55	5.91	5.26	3.50	4.13	5.08	5.08			
	OVARC1000254	16.05	11.01	13.12	50.15	59.76	29.83	42.38	33.82	33.82	*	+	**
30	OVARC1000255	5.5	3.14	2.99	5.45	4.17	3.19	3.91	4.30	4.3			
	OVARC1000267	8.95	5.90	5.53	9.61	7.91	10.70	8.96	10.59	10.59			
	OVARC1000275	0.38	0.28	0.65	1.7	1.69	1.90	10.31	9.09	9.09	**	+	**
	OVARC1000287	2.16	1.07	1.61	5.38	6.97	4.90	26.09	33.14	33.14	**	+	**
	OVARC1000288	7.99	3.43	4.43	6.36	6.18	3.91	4.34	4.81	4.81			
35	OVARC1000298	8.86	6.47	4.36	11.32	12.55	7.25	6.14	7.12	7.12			
	OVARC1000302	3.96	1.75	1.50	3.75	4.71	3.28	2.04	3.19	3.19			
	OVARC1000304	6.08	4.82	3.98	7.97	7.57	5.26	4.58	6.93	6.93			
	OVARC1000307	5.1	1.95	3.30	4.25	2.68	4.18	3.69	3.54	3.54			
	OVARC1000309	6.17	3.11	3.95	6.94	5.55	4.98	5.49	5.61	5.61			
40	OVARC1000312	4.47	2.31	2.62	3.43	3.39	3.03	5.14	4.44	4.44			
	OVARC1000313	7.23	3.04	5.41	6.92	6.31	4.37	7.31	10.70	10.7			
	OVARC1000321	8.81	5.88	6.66	13.97	15.87	13.56	14.26	12.53	12.53	**	+	**
	OVARC1000326	3.94	3.57	2.28	3.59	3.18	3.94	3.62	3.71	3.71			
	OVARC1000327	4.66	2.13	3.59	7.38	4.82	4.34	3.97	5.68	5.68			
	OVARC1000331	6.82	4.80	4.04	7.15	6.72	8.39	4.61	6.40	6.4			
45	OVARC1000335	5.22	3.45	3.68	6.19	5.78	6.01	4.99	5.32	5.32	*	+	
	OVARC1000347	2.86	2.21	1.39	1.74	2.06	3.33	1.79	3.03	3.03			
	OVARC1000348	7.01	4.29	4.68	13.43	12.42	16.47	7.65	8.17	8.17	**	+	*
	OVARC1000363	4.22	3.97	3.08	6.15	6.28	7.74	2.83	4.38	4.38	**	+	
	OVARC1000377	2.82	1.76	1.53	3.08	2.53	1.71	0.35	2.23	2.23			
50	OVARC1000382	5.76	1.98	3.91	4.79	5.06	3.60	4.10	6.90	6.9			
	OVARC1000384	6.02	5.30	4.11	6.76	8.20	10.33	8.85	9.44	9.44	*	+	**
	OVARC1000401	2.8	1.75	1.96	2.86	2.28	2.89	2.67	3.48	3.48			
	OVARC1000406	114.78	80.20	88.37	88.01	73.54	119.34	90.62	95.77	95.77			
	OVARC1000407	4.6	3.44	3.17	5.58	5.61	8.84	5.08	4.38	4.38			
55	OVARC1000408	16.3	13.53	12.64	45.51	42.22	49.78	26.85	32.12	32.12	**	+	**
	OVARC1000410	6.71	4.55	5.34	4.6	7.22	6.74	6.55	6.11	6.11			
	OVARC1000411	3.32	1.84	2.60	3.91	7.15	2.94	1.78	2.39	2.39			

Table 292

	NT2RP4002894	30.47	15.42	16.30	15.33	13.44	13.63	14.61	7.84	11.34			
5	NT2RP4002896	5.01	2.57	1.03	5.77	4.90	3.35	4.85	5.20	6.5			
	NT2RP4002905	3.65	2.18	-2.47	3.73	2.63	3.46	2.67	3.64	2.22			
	NT2RP4002907	6.79	1.23	2.84	16.01	14.42	10.02	12.06	10.10	6.54	*	+	
	NT2RP5003459	65.35	36.44	48.17	27.67	30.09	25.05	9.64	20.91	22.09		*	-
	NT2RP5003461	4.58	3.60	3.17	6.87	4.80	7.46	3.05	4.17	2.86	*	+	
10	NT2RP5003471	5.96	3.26	3.68	5.59	5.78	6.38	36.49	36.45	36.48		**	+
	NT2RP5003477	4.19	2.26	3.16	4.58	5.06	6.58	6.46	4.38	3.4			
	NT2RP5003487	220.55	93.22	98.28	181.60	187.80	154.38	86.45	85.87	93.23			
	NT2RP5003492	7.41	4.46	3.61	6.80	6.09	7.24	6.01	5.67	4.83			
	NT2RP5003500	3.73	2.01	1.80	4.33	3.62	5.68	2.91	3.93	3.14			
15	NT2RP5003506	9.63	4.24	5.17	6.58	8.38	7.49	5.4	7.54	7.66			
	NT2RP5003512	2.05	1.82	0.90	1.93	2.76	1.89	1.76	3.04	2.68			
	NT2RP5003522	5.00	3.31	4.09	6.05	5.02	4.70	4.69	4.96	3.11			
	NT2RP5003524	2.66	1.03	1.85	3.05	3.14	2.14	2.01	1.80	0.86			
	NT2RP5003527	27.32	17.39	28.11	33.15	29.19	33.15	34.18	28.33	30.99	*	+	*
20	NT2RP5003531	6.09	4.05	3.52	14.63	15.87	11.17	18.91	10.15	13.33	**	+	*
	NT2RP5003534	4.69	3.24	2.48	4.56	5.46	3.21	3.85	3.74	4.1			
	NT2RP6000020	14.93	5.50	7.94	19.43	12.24	14.47	28	17.69	22.01		*	+
	NT2RP6000022	2.09	1.92	1.10	2.89	3.69	3.48	1.85	3.95	3.04	*	+	
	NT2RP6000050	6.72	2.85	2.69	5.15	4.13	6.91	3.13	4.74	4.15			
25	NT2RP6000063	4.32	1.86	2.74	4.12	3.95	5.49	4.77	5.84	5.17		*	+
	NT2RP6000074	7.65	3.63	3.82	5.82	4.62	5.47	3.91	5.25	4.12			
	NT2RP6000083	7.65	4.46	4.22	5.62	7.05	9.12	4.96	6.80	6.49			
	NT2RP6000100	8.20	3.69	3.69	11.31	10.03	10.20	5.69	6.11	4.22	*	+	
	NT2RP6000123	8.42	4.03	3.87	7.40	6.54	4.76	5.08	5.14	4.33			
	NT2RP6000129	5.14	2.45	3.11	3.95	4.30	4.21	3.96	4.16	4.57			
30	NT2RP6000147	3.79	2.50	3.26	15.24	15.27	11.86	26.48	14.22	25.1	**	+	**
	NT2RP6000163	1.43	1.14	1.15	3.25	1.30	2.00	1.02	2.54	1.73			
	NT2RP6000181	7.19	4.67	4.25	6.16	6.80	4.73	6.67	5.10	6.2			
	NT2RP6000182	5.25	3.12	3.43	5.76	4.23	7.79	3.45	3.70	2.44			
35	OVARC1000001	4.47	2.05	2.92	5.01	4.27	3.71	5.92	4.78	4.37			
	OVARC1000003	4.03	2.27	2.17	3.53	4.26	1.98	1.87	2.81	4.16			
	OVARC1000004	69.94	45.81	40.28	31.28	33.52	34.13	14.2	20.99	22.91		*	-
	OVARC1000006	2.75	1.60	1.91	3.55	3.17	2.27	3.59	3.71	3.52		*	+
	OVARC1000013	3.58	2.31	1.87	3.88	4.15	3.20	3.52	4.55	2.95			
	OVARC1000014	5.72	2.95	3.69	6.24	6.32	5.61	4.07	4.99	4.34			
40	OVARC1000017	6.14	3.05	3.33	4.90	5.12	5.05	3.15	5.17	5.31			
	OVARC1000026	55.69	36.49	45.68	51.02	60.13	48.46	28.42	36.95	25.22			
	OVARC1000035	9.77	8.46	8.93	13.12	14.00	9.30	7.02	5.89	5.3		**	-
	OVARC1000037	31.27	16.99	12.47	49.92	39.93	32.59	18.22	25.08	32.08			
	OVARC1000058	10.77	5.52	3.11	12.87	13.32	13.63	6.74	5.82	8.66	*	+	
	OVARC1000060	3.24	1.54	1.26	3.04	2.70	2.45	2.09	2.66	3.05			
45	OVARC1000068	2.38	1.15	1.10	3.07	2.77	1.87	1.01	3.23	1.66			
	OVARC1000069	4.64	2.24	2.58	7.95	8.04	5.29	4.94	7.33	5.21	*	+	
	OVARC1000071	4.18	2.24	2.19	3.21	4.19	2.93	1.32	4.38	1.25			
	OVARC1000075	116.66	59.06	70.03	104.67	109.44	102.05	127.1	180.67	194.9		*	+
	OVARC1000083	16.13	9.03	10.85	16.27	15.52	17.85	9.32	13.62	11.15			
50	OVARC1000085	90.31	52.35	57.44	84.93	91.25	74.75	46.89	55.51	55.51			
	OVARC1000086	3.63	2.07	4.18	7.09	7.77	8.13	5.87	6.77	6.77	**	+	*
	OVARC1000087	2.46	0.70	0.93	1.65	1.80	2.44	2.22	3.58	3.58			
	OVARC1000090	7.22	4.69	6.24	15.64	14.18	15.90	5.67	9.11	9.11	**	+	
	OVARC1000091	3.66	1.42	2.09	5.7	4.57	5.66	4.01	3.77	3.77	*	+	
55	OVARC1000092	3.91	1.98	2.18	6.09	6.36	8.26	4.35	4.86	4.86	*	+	*
	OVARC1000105	11.95	8.25	9.35	12.3	11.58	13.87	6.66	8.05	8.05			
	OVARC1000106	23.29	10.32	10.91	20.75	17.39	12.69	12.13	18.29	18.29			

Table 291

	NT2RP4002186	24.35	16.62	12.92	73.40	68.88	51.66	18.99	20.53	42.77	**	+		
5	NT2RP4002187	16.88	9.15	8.08	13.25	18.23	17.99	14.62	13.98	23.37				
	NT2RP4002188	9.49	5.18	4.64	14.32	14.99	9.78	4.92	6.78	9.43	*	+		
	NT2RP4002199	3.33	0.85	1.71	2.01	2.76	1.40	1.46	4.34	2.92				
	NT2RP4002206	7.79	3.61	3.56	5.56	5.23	3.75	3.53	5.24	4.66				
	NT2RP4002210	3.95	1.94	2.05	3.42	2.86	2.32	2.13	4.76	2.28				
10	NT2RP4002222	4.87	2.50	3.89	4.48	5.59	3.24	4.1	4.89	3.82				
	NT2RP4002241	10.39	8.75	9.34	8.11	10.75	7.80	3.37	5.39	6.12		**	-	
	NT2RP4002248	5.75	3.15	2.68	4.58	3.49	3.31	6.08	4.55	3.57				
	NT2RP4002250	2.77	1.28	0.36	1.28	1.49	1.07	2.02	0.58	1.13				
	NT2RP4002259	11.44	4.70	6.93	10.37	10.26	7.96	6.18	7.00	6.72				
15	NT2RP4002268	9.49	7.15	6.70	7.16	8.97	8.79	12.35	10.44	12.35		*	+	
	NT2RP4002288	23.22	15.06	19.08	20.88	28.68	23.53	20.32	17.25	20.1				
	NT2RP4002290	9.48	5.25	5.05	15.46	15.55	18.46	13.55	11.18	12.37	**	+	*	+
	NT2RP4002298	5.94	3.63	4.51	10.11	6.35	12.09	3.11	5.17	4.75				
	NT2RP4002306	5.29	2.43	3.39	8.59	7.82	9.25	3.86	4.05	3.61	**	+		
20	NT2RP4002308	2.50	1.35	1.43	1.70	2.93	1.47	2.72	1.97	2.14				
	NT2RP4002336	9.03	4.10	4.50	6.72	4.54	7.26	5.89	4.31	4.91				
	NT2RP4002340	0.95	0.34	0.60	0.63	0.88	0.24	1.51	1.53	0.76				
	NT2RP4002361	3.28	2.38	1.78	3.90	2.34	2.47	2.23	2.16	1.92				
	NT2RP4002367	3.30	2.19	1.54	3.77	4.95	3.32	2.84	2.25	3				
	NT2RP4002368	4.21	2.40	3.66	5.83	4.14	3.92	5.91	4.62	3.42				
25	NT2RP4002377	3.62	4.26	2.84	5.85	2.38	5.20	4.75	3.54	3.33				
	NT2RP4002408	29.46	20.49	24.43	3.81	2.37	2.48	1.32	0.66	1.06	**	-	**	-
	NT2RP4002425	1.74	1.67	0.75	1.77	1.60	1.39	2.92	1.48	1.25				
	NT2RP4002432	8.35	5.60	3.82	5.76	5.85	4.41	8.08	6.14	6.6				
	NT2RP4002447	9.10	3.90	3.22	12.78	11.88	10.40	5.91	5.47	6.48	*	+		
30	NT2RP4002451	2.21	2.30	1.71	3.91	4.29	3.31	5.98	6.15	6.01	**	+	**	+
	NT2RP4002461	7.09	5.26	5.72	12.39	9.75	9.13	7.77	8.39	7.06	*	+		
	NT2RP4002486	5.84	4.56	5.50	5.14	5.35	4.72	7.44	6.30	6.54		*	+	
	NT2RP4002517	3.21	2.30	2.48	3.27	2.89	3.72	3.06	3.47	2.44				
	NT2RP4002556	10.73	5.00	5.11	11.36	8.97	7.80	4.9	4.20	4.51				
35	NT2RP4002569	5.60	3.78	2.56	4.11	4.44	3.67	5.29	5.70	3.72				
	NT2RP4002587	2.41	1.81	1.87	2.59	3.67	3.36	7.6	6.60	7.95	*	+	**	+
	NT2RP4002591	7.42	6.05	5.29	12.68	12.07	10.38	7.78	4.95	7.6	**	+		
	NT2RP4002607	6.11	2.67	2.59	6.08	4.47	5.73	3.49	4.31	2.91				
	NT2RP4002627	5.30	4.31	4.08	5.45	8.00	6.98	9.55	7.80	7.44		**	+	
40	NT2RP4002628	13.62	7.50	7.90	12.59	11.82	9.24	5.81	7.23	4.46				
	NT2RP4002630	3.81	2.47	2.90	6.00	2.15	4.82	6.13	6.96	4.18		*	+	
	NT2RP4002639	4.77	2.18	3.85	2.27	2.26	2.48	1.79	3.34	1.18				
	NT2RP4002641	8.72	3.54	3.33	4.53	5.23	4.41	5.45	5.35	8.22				
	NT2RP4002658	39.52	16.53	21.90	10.69	10.22	8.63	12.92	12.25	13.09				
	NT2RP4002669	8.68	5.48	3.61	6.49	4.90	5.66	4.3	4.66	5.21				
45	NT2RP4002677	11.90	7.10	10.78	11.62	13.84	10.32	4.5	5.32	4.63		*	-	
	NT2RP4002715	6.49	4.85	5.45	16.06	11.33	12.78	11.52	13.89	13.89	**	+	**	+
	NT2RP4002750	11.19	4.76	5.94	4.82	3.94	4.94	3.86	3.11	3.58				
	NT2RP4002784	5.22	3.74	4.33	6.90	5.66	7.76	7.61	4.05	3.39	*	+		
	NT2RP4002791	2.32	2.02	2.01	4.89	4.01	4.73	3.62	3.85	2.33	**	+		
50	NT2RP4002811	6.07	3.91	2.96	1.95	3.45	3.30	4.41	5.24	4.51				
	NT2RP4002830	11.00	4.98	5.60	10.88	8.44	6.08	6.46	6.96	4.35				
	NT2RP4002832	2.65	2.09	2.28	3.38	2.40	2.76	1.95	3.23	1.27				
	NT2RP4002850	10.22	7.64	6.28	14.24	10.59	11.13	9.28	9.04	5.75				
	NT2RP4002874	3.50	2.69	1.87	3.65	3.03	4.18	3.78	5.05	2.68				
55	NT2RP4002884	17.66	6.25	9.46	10.83	9.85	8.92	15.05	14.77	10.31				
	NT2RP4002888	20.83	12.71	14.10	15.29	12.54	11.78	19.91	18.79	15.14				
	NT2RP4002891	6.49	3.33	5.04	17.64	15.92	12.46	8.11	7.56	7.35	**	+	*	+

Table 290

	NT2RP4001841	5.15	2.19	2.44	6.33	5.75	3.95	4.94	4.03	3.03				
5	NT2RP4001849	4.08	2.37	1.90	1.96	2.08	2.74	2.12	3.59	2.22				
	NT2RP4001861	19.55	11.05	8.48	18.06	19.21	17.61	12.49	10.31	10.34				
	NT2RP4001877	18.38	12.98	11.71	13.65	17.92	15.26	10.17	11.03	9.86				
	NT2RP4001879	6.00	4.86	5.20	4.62	6.88	7.55	4.96	6.52	5.75				
	NT2RP4001889	3.83	2.48	2.26	4.36	5.15	5.12	3.39	5.09	3.84	*	+		
10	NT2RP4001893	4.85	2.58	3.31	5.78	4.46	6.55	5.02	4.75	1.96				
	NT2RP4001896	4.86	2.86	3.13	4.46	5.44	4.95	3.44	3.93	1.91				
	NT2RP4001898	12.63	7.18	6.38	11.85	13.48	14.72	8.27	7.05	8.92				
	NT2RP4001901	9.37	5.10	4.58	7.22	7.41	7.58	5.92	5.84	4.25				
	NT2RP4001910	44.22	14.42	25.27	36.18	28.56	31.03	15.44	16.11	13.43				
15	NT2RP4001925	6.01	3.53	4.07	7.13	8.88	6.52	5.38	5.68	3.89	*	+		
	NT2RP4001926	5.02	2.32	4.10	6.70	3.01	7.01	3.35	4.83	1.34				
	NT2RP4001927	7.81	3.22	8.37	2.90	3.77	4.75	2.11	3.46	2.61				
	NT2RP4001931	12.13	7.10	9.23	9.30	11.80	10.57	7.09	9.58	5.89				
	NT2RP4001933	7.27	5.93	8.24	33.37	26.48	21.53	12.07	15.48	9.59	**	+	*	+
20	NT2RP4001938	11.79	6.36	5.51	7.00	8.59	7.23	7.68	7.54	9.66				
	NT2RP4001942	19.13	10.55	10.00	11.76	13.07	12.47	8.35	7.90	8.71				
	NT2RP4001945	3.39	2.16	1.75	1.10	2.83	1.75	3.88	3.65	3.03				
	NT2RP4001946	2.78	2.76	2.10	6.68	5.62	8.03	3.2	4.28	3.28	**	+		
	NT2RP4001947	0.70	0.50	0.71	3.55	3.12	4.05	1.69	2.42	0.29	**	+		
25	NT2RP4001950	52.07	29.14	30.34	3.90	3.31	3.63	2.85	3.53	3.23	*	-	*	-
	NT2RP4001953	6.50	3.60	5.67	12.09	12.07	9.95	5.86	6.12	3.31	**	+		
	NT2RP4001966	3.87	2.06	1.81	2.93	2.33	3.06	2.56	3.55	1.61				
	NT2RP4001970	18.77	7.73	6.33	7.39	9.12	8.12	6.83	7.05	6.87				
	NT2RP4001975	16.12	8.35	8.50	16.73	14.58	16.13	21.64	17.08	14.87				
	NT2RP4001988	6.11	2.52	2.36	2.17	2.97	2.42	4.05	5.29	6.8				
30	NT2RP4001996	8.88	6.41	7.06	5.35	6.06	5.33	4.86	5.72	5.5				
	NT2RP4002014	5.46	3.70	3.51	5.82	4.28	3.92	5.71	6.94	6.45		*	+	
	NT2RP4002018	4.51	3.12	2.83	6.79	4.88	5.98	4.66	10.23	5.14	*	+		
	NT2RP4002035	6.12	4.46	6.67	7.19	6.57	6.76	5.8	5.73	6.32				
	NT2RP4002043	17.40	10.99	15.66	15.62	10.19	12.89	8.93	9.28	8.15		*	-	
35	NT2RP4002046	6.17	4.77	3.90	3.50	9.38	4.20	6.26	5.07	7.72				
	NT2RP4002047	14.83	7.78	9.72	12.74	11.88	9.86	4.4	4.88	5.22		*	-	
	NT2RP4002052	3.82	2.22	2.36	3.72	2.89	4.12	4.34	4.20	5.03		*	+	
	NT2RP4002056	55.72	38.98	47.46	51.12	52.01	41.19	44.9	38.97	37.38				
	NT2RP4002057	17.74	8.34	10.35	10.25	6.84	10.23	9.46	9.43	9				
40	NT2RP4002058	5.05	3.72	3.60	3.34	2.84	3.35	3.74	3.86	2.96				
	NT2RP4002064	2.43	1.64	1.15	2.53	2.72	2.44	2.13	3.96	2.74				
	NT2RP4002071	6.91	5.83	6.59	9.94	11.45	10.50	6.83	7.79	5.44	**	+		
	NT2RP4002075	5.65	2.21	2.77	1.76	1.64	2.01	1.03	0.80	1.27				
	NT2RP4002078	12.20	5.57	6.28	21.16	11.84	9.58	9.34	5.65	7.44				
	NT2RP4002081	8.20	4.41	4.38	8.71	5.52	5.98	8.56	5.96	6.86				
45	NT2RP4002083	1.41	0.64	0.77	1.12	0.92	0.88	1.16	1.92	2.62				
	NT2RP4002099	3.50	1.74	2.24	2.98	2.94	2.77	2.69	3.45	2.97				
	NT2RP4002106	16.08	11.97	16.65	14.50	11.42	13.37	8.7	8.53	7.16		*	-	
	NT2RP4002111	14.95	7.66	10.77	13.64	14.70	14.45	16.75	17.55	15.81				
	NT2RP4002112	5.99	2.81	3.54	4.57	5.85	6.57	6.12	5.13	4.94				
50	NT2RP4002116	14.14	7.04	5.48	14.30	12.58	11.93	8.14	4.91	6.09				
	NT2RP4002122	15.83	9.46	8.25	6.72	6.57	5.28	1.63	2.27	1.64		*	-	
	NT2RP4002126	7.11	2.89	3.58	2.17	3.83	2.41	4.77	4.31	5.35				
	NT2RP4002133	10.15	4.28	5.52	6.19	8.16	5.73	5.02	6.44	4.79				
	NT2RP4002136	13.83	8.55	8.39	5.76	6.14	5.01	5.63	5.28	4.82		*	-	
55	NT2RP4002139	25.38	27.01	30.04	25.35	29.41	24.98	13.23	24.52	19.91				
	NT2RP4002174	3.31	1.15	2.46	3.71	3.16	4.15	3.14	5.41	4.47				
	NT2RP4002185	10.77	7.55	7.67	15.20	13.59	13.41	10.77	8.24	8.8	*	+		

Table 289

	NT2RP4001529	9.24	4.27	3.42	3.66	4.21	3.95	6.65	3.78	5.28				
	NT2RP4001531	7.58	4.22	3.87	4.40	6.79	5.07	4.85	4.25	5.33				
5	NT2RP4001546	27.96	14.34	13.14	33.50	26.35	22.36	39.72	37.62	23.88				
	NT2RP4001547	5.16	3.87	3.59	6.27	5.81	5.41	6.77	5.69	7.74	*	+	*	+
	NT2RP4001551	4.66	2.25	2.91	1.72	2.50	2.23	1.06	2.31	2.02				
	NT2RP4001555	2.63	1.70	1.48	1.84	1.34	1.78	3.29	2.29	1.99				
10	NT2RP4001567	4.17	2.21	3.48	5.17	4.12	2.97	3.53	3.55	4.6				
	NT2RP4001568	24.66	11.55	19.71	26.48	16.71	27.97	21.61	20.91	21.83				
	NT2RP4001569	13.23	7.51	6.17	8.88	7.94	7.65	6.86	6.56	7.44				
	NT2RP4001571	3.88	2.14	1.80	4.74	3.69	4.71	3.97	5.20	7.86				
	NT2RP4001574	8.96	4.84	4.26	8.19	9.78	5.65	6.26	6.22	8.16				
15	NT2RP4001575	8.04	4.77	3.76	6.08	7.50	5.82	4.63	5.56	5.85				
	NT2RP4001578	11.18	4.73	6.33	7.50	4.87	4.81	7.41	8.00	7.35				
	NT2RP4001592	9.35	5.87	4.90	5.95	6.70	4.56	3.37	8.97	5.41				
	NT2RP4001593	6.28	4.83	5.72	9.71	12.44	12.90	7.66	7.56	6.44	**	+	*	+
	NT2RP4001605	4.40	2.61	3.07	7.26	7.76	5.64	5.16	7.35	8.18	*	+	*	+
20	NT2RP4001606	13.15	5.10	4.06	9.17	7.65	6.75	3.7	4.31	6.28				
	NT2RP4001607	3.47	1.57	1.29	3.76	4.78	2.65	1.67	3.06	4.34				
	NT2RP4001610	4.08	2.08	1.47	3.77	3.73	2.68	2.34	4.35	2.92				
	NT2RP4001614	2.75	1.07	1.10	2.96	1.97	1.29	2.18	3.56	3.15				
	NT2RP4001623	3.08	1.60	1.52	2.58	2.94	2.80	1.24	3.23	2.34				
	NT2RP4001626	19.42	15.83	18.19	15.38	17.59	13.04	1.75	4.18	2.95		**	-	
25	NT2RP4001634	4.38	2.77	2.43	4.92	4.36	4.52	1.82	3.51	2.53				
	NT2RP4001638	2.68	1.70	0.84	1.98	2.75	2.80	1.64	3.48	1.26				
	NT2RP4001644	3.61	2.50	2.30	4.35	3.54	2.45	4.35	2.84	4.05				
	NT2RP4001646	20.39	11.21	10.21	30.98	19.98	25.17	21.75	14.88	9.56				
	NT2RP4001656	6.55	3.72	4.64	5.20	5.23	4.49	4.29	3.23	2.79				
30	NT2RP4001666	5.11	3.28	3.35	4.54	4.56	3.95	3.53	3.52	3.5				
	NT2RP4001670	7.31	3.77	5.28	4.59	6.96	4.67	4.23	4.15	4.55				
	NT2RP4001677	16.68	12.12	14.19	29.06	40.57	32.81	33.86	36.13	36.39	**	+	**	+
	NT2RP4001679	11.61	4.52	5.94	19.33	14.25	14.99	8.64	9.90	7.91	*	+		
	NT2RP4001695	20.41	7.98	11.64	19.72	19.63	15.23	7.89	9.75	7.32				
35	NT2RP4001696	6.64	4.27	3.64	4.33	3.58	5.85	4.75	3.99	3.79				
	NT2RP4001699	1.63	1.58	0.71	2.91	1.63	2.15	3.74	2.30	2.42				
	NT2RP4001717	5.33	4.49	3.61	5.92	6.26	5.39	5.73	6.49	5.79				
	NT2RP4001719	3.81	3.40	2.34	4.26	2.94	3.04	4.14	3.43	2.54				
	NT2RP4001725	4.09	3.08	1.88	3.37	4.40	3.86	2.62	4.74	3.15				
	NT2RP4001726	4.90	3.18	3.91	4.82	4.39	4.14	4.14	5.24	5.01				
40	NT2RP4001730	0.78	0.69	0.71	1.42	1.12	2.01	0.61	1.16	0.59	*	+		
	NT2RP4001739	4.83	2.71	3.87	5.22	3.09	4.63	4.39	5.41	4.57				
	NT2RP4001741	10.82	7.34	4.37	12.44	9.41	10.54	7.99	6.39	5.79				
	NT2RP4001753	11.73	4.55	5.91	14.42	16.38	12.01	9.64	6.92	8.76				
	NT2RP4001760	12.48	7.93	6.76	6.21	7.16	7.70	2.14	2.05	1.86		*	-	
45	NT2RP4001787	45.15	35.87	34.25	46.58	51.77	52.14	19.69	24.09	19.41	*	+	**	-
	NT2RP4001790	6.06	3.59	2.88	5.91	6.74	7.08	5.5	5.13	5.27				
	NT2RP4001795	25.43	15.84	22.47	18.33	17.56	16.99	11.05	11.79	10.2		*	-	
	NT2RP4001803	3.51	2.55	1.55	5.77	4.36	4.15	4.43	4.25	2.97	*	+		
	NT2RP4001805	4.04	2.46	2.43	5.53	4.54	4.71	3.91	2.59	3.66	*	+		
50	NT2RP4001809	14.99	9.07	7.27	11.92	10.72	9.25	11.36	11.25	11.16				
	NT2RP4001817	16.10	8.59	7.80	8.81	9.92	9.75	5.74	6.19	5.7				
	NT2RP4001822	9.90	6.09	4.79	7.82	4.55	6.51	6.73	5.44	6.61				
	NT2RP4001823	1.63	1.96	0.82	1.62	2.17	1.74	1.56	1.67	0.88				
	NT2RP4001827	5.09	4.68	4.45	4.54	5.32	5.79	7.53	6.64	8.76		*	+	
	NT2RP4001828	17.04	10.89	10.46	15.89	15.47	13.14	13.38	12.00	9.76				
55	NT2RP4001836	5.07	3.08	3.80	4.72	5.04	5.75	5.07	4.56	2.8				
	NT2RP4001838	6.83	3.89	5.07	5.21	5.01	6.41	4.27	6.56	2.85				

Table 288

5	NT2RP4001228	6.93	2.54	3.03	5.28	9.41	5.96	8.24	4.93	8.37				
	NT2RP4001235	6.11	4.31	3.21	5.70	5.94	5.25	5.94	4.41	5.1				
	NT2RP4001256	4.51	1.77	2.22	4.07	5.11	4.94	4.27	3.05	2.43				
	NT2RP4001257	6.40	4.02	2.26	5.05	5.54	3.44	5.95	5.21	4.31				
	NT2RP4001260	5.39	3.07	4.18	8.97	9.59	5.62	5.8	6.24	6.64		*	+	
	NT2RP4001261	14.65	12.44	12.58	14.19	12.55	13.99	17.34	12.10	15.2				
10	NT2RP4001274	4.71	4.57	4.07	7.45	6.65	6.76	5.26	6.13	6.26	**	+	*	+
	NT2RP4001276	15.31	8.46	8.50	10.61	14.38	10.37	11.44	11.39	8.98				
	NT2RP4001283	63.21	34.01	32.33	24.21	25.03	19.31	48.06	42.63	46.56				
	NT2RP4001299	15.00	9.02	6.78	6.64	8.24	7.13	7.92	6.14	6.14				
	NT2RP4001313	3.06	1.56	1.37	2.51	0.89	2.21	1.62	2.23	2.1				
15	NT2RP4001315	3.67	2.67	2.40	3.95	5.09	3.45	3.89	3.89	4.16				
	NT2RP4001320	9.02	4.65	5.15	9.20	8.51	8.68	15.43	12.65	14.49		**	+	
	NT2RP4001325	12.74	11.37	11.78	16.64	15.36	9.87	12.12	10.53	7.42				
	NT2RP4001336	6.40	4.16	5.13	5.38	3.83	5.19	4.39	4.05	2.52				
	NT2RP4001339	3.62	2.24	4.32	4.37	4.09	4.92	3.51	4.78	3.43				
20	NT2RP4001343	8.44	4.63	3.67	7.94	6.79	5.81	5.7	6.09	6.51				
	NT2RP4001344	5.76	3.40	4.09	5.03	5.50	6.54	6.12	6.22	5.58				
	NT2RP4001345	6.21	3.12	2.61	3.29	6.07	5.15	4.25	4.33	4.38				
	NT2RP4001351	11.92	6.04	5.53	9.86	6.47	8.71	6.54	7.28	6.61				
	NT2RP4001353	1.80	1.08	1.42	2.16	2.00	2.04	2.15	2.48	2.23	*	+	*	+
	NT2RP4001355	2.54	1.08	2.05	2.40	2.01	1.99	2.51	3.62	2.23				
25	NT2RP4001367	23.22	13.41	17.84	6.30	4.94	5.47	9.28	11.30	7.57	*	-	*	-
	NT2RP4001372	5.35	2.77	2.56	3.34	4.53	3.59	4.57	5.24	5.57				
	NT2RP4001373	10.60	5.25	4.77	8.11	9.86	9.53	6.1	5.34	6.98				
	NT2RP4001375	5.11	3.33	2.60	2.66	4.56	3.81	2.85	3.42	3.31				
	NT2RP4001379	3.86	2.14	2.09	2.83	2.70	4.72	3.26	3.43	2.58				
30	NT2RP4001381	8.37	5.24	5.75	10.66	11.10	10.55	6.09	7.62	6.54	*	+		
	NT2RP4001386	3.36	2.18	2.25	6.41	4.78	6.49	3.68	5.89	3.24	**	+		
	NT2RP4001389	10.33	5.90	8.63	13.74	8.10	10.59	13.58	10.92	11.95				
	NT2RP4001396	1.51	0.17	0.39	1.10	1.45	1.19	1.43	2.48	0.52				
	NT2RP4001407	2.74	1.02	1.62	3.87	3.78	1.98	2.72	2.67	1.52				
35	NT2RP4001409	7.90	3.42	3.68	8.04	5.25	6.08	3.89	2.35	3.87				
	NT2RP4001410	41.71	16.67	20.24	29.88	31.04	31.69	28.88	20.00	22.74				
	NT2RP4001414	11.73	6.50	5.48	10.69	11.38	10.17	10.68	8.69	10.89				
	NT2RP4001424	3.25	2.51	1.43	4.18	3.70	4.01	2.5	5.15	3.66	*	+		
	NT2RP4001433	10.93	1.50	1.13	15.16	15.56	3.13	10.41	4.52	7				
40	NT2RP4001438	8.06	6.23	6.43	14.12	10.57	11.39	6.77	9.65	7.69	*	+		
	NT2RP4001442	5.25	2.76	3.72	6.62	2.55	2.88	2.74	3.33	2.46				
	NT2RP4001447	1.94	1.07	2.00	4.12	2.36	3.98	1.68	3.22	0.71	*	+		
	NT2RP4001466	13.13	5.79	4.82	7.69	5.30	6.70	2.91	4.53	3.9				
	NT2RP4001467	4.50	1.22	1.33	0.82	1.55	1.40	3.66	4.13	3.7				
	NT2RP4001472	4.77	3.08	3.33	7.29	7.84	10.23	7.79	8.21	9.21	*	+	**	+
45	NT2RP4001474	2.86	1.72	1.90	2.18	3.93	2.05	1.94	3.80	3.06				
	NT2RP4001483	2.29	1.49	1.84	3.04	2.50	2.14	2.24	3.68	2.54				
	NT2RP4001488	5.16	2.65	2.75	5.33	5.10	5.16	4.15	4.07	6.19				
	NT2RP4001492	5.93	3.30	2.87	5.58	3.40	4.66	3.78	4.60	5.29				
	NT2RP4001498	2.17	1.63	1.33	2.59	1.07	2.19	2.61	1.92	1.74				
50	NT2RP4001502	36.00	12.08	15.43	15.15	11.96	14.06	11.15	10.06	10.33				
	NT2RP4001503	12.74	6.75	6.97	11.88	9.69	8.87	5.71	4.79	6.02				
	NT2RP4001507	5.29	3.55	4.09	6.91	8.58	5.74	3.85	4.47	6.06	*	+		
	NT2RP4001510	9.01	6.05	7.69	15.28	11.96	14.90	7.45	6.33	7.03	**	+		
	NT2RP4001516	6.51	3.15	3.51	3.57	3.42	3.81	3.63	4.76	4.46				
55	NT2RP4001520	26.12	11.82	16.11	17.96	13.99	17.71	16.8	15.23	12.31				
	NT2RP4001523	3.37	1.82	2.58	4.23	3.77	4.26	2.29	4.88	4.21	*	+		
	NT2RP4001524	11.16	7.76	6.79	8.80	7.75	9.91	6.38	9.28	5.14				

Table 287

5	NT2RP4000990	0.91	1.17	0.68	5.32	4.83	4.20	3.51	3.92	3.51	**	+	**	+
	NT2RP4000994	6.03	3.61	2.39	2.73	3.58	3.95	4.94	3.50	5.8				
	NT2RP4000996	6.29	4.22	3.37	8.35	8.21	4.36	4.41	5.02	6.24				
	NT2RP4000997	61.78	21.49	33.43	48.43	44.30	38.85	25.67	23.78	20.69				
	NT2RP4001001	5.72	4.90	3.47	5.67	6.31	7.83	5.36	5.68	6.44				
	NT2RP4001004	2.47	1.20	1.29	1.66	1.42	2.31	0.88	2.30	2.26				
10	NT2RP4001006	6.01	3.42	6.46	5.11	3.94	7.35	4.19	4.92	4.66				
	NT2RP4001009	8.55	4.50	6.33	9.69	4.66	6.57	7.89	8.50	7.3				
	NT2RP4001010	2.33	1.99	3.31	3.50	2.89	4.49	3.41	2.18	2.22				
	NT2RP4001013	24.76	12.16	10.77	11.37	8.47	9.68	9.97	7.96	8.93				
	NT2RP4001029	12.87	4.18	5.93	5.61	5.98	4.93	3.75	3.75	3.77				
15	NT2RP4001036	12.25	7.10	7.56	11.16	11.59	9.83	8.7	8.94	6.61				
	NT2RP4001041	12.91	6.26	9.00	10.06	7.34	6.55	5.46	5.13	5.26				
	NT2RP4001042	19.25	12.69	10.60	14.77	15.99	12.64	7.69	8.09	6.86				
	NT2RP4001046	7.12	4.49	4.11	7.61	7.51	8.81	6.32	5.77	5.49				
	NT2RP4001050	2.62	1.51	1.21	2.43	2.08	3.36	1.88	2.76	1.6				
20	NT2RP4001051	6.34	2.77	3.34	9.61	5.53	9.29	3.29	7.15	4.7				
	NT2RP4001057	8.53	5.25	3.91	5.29	5.25	3.31	3.9	2.95	5.29				
	NT2RP4001063	10.42	5.01	5.86	6.23	5.90	5.66	7.16	6.14	5.17				
	NT2RP4001064	8.38	3.24	3.12	6.83	5.16	4.26	7.84	6.40	8.7				
	NT2RP4001067	3.31	1.58	2.32	2.67	2.92	2.43	3.23	3.73	3.54				
	NT2RP4001078	4.41	2.35	1.43	3.09	2.15	3.81	2.07	3.77	2.39				
25	NT2RP4001079	3.33	2.47	3.40	5.24	5.12	4.58	4.51	5.49	6.01	**	+	*	+
	NT2RP4001080	1.87	1.13	0.72	2.09	1.21	1.64	1.8	2.65	1.93				
	NT2RP4001086	6.48	4.50	4.95	6.91	6.12	6.66	5.1	5.61	4.86				
	NT2RP4001095	9.39	3.28	2.95	11.12	8.02	6.83	6.4	5.11	6.99				
	NT2RP4001098	8.66	3.42	3.13	5.99	6.59	3.50	4.06	3.58	3.83				
30	NT2RP4001100	15.58	6.86	5.99	15.36	16.25	10.53	11.07	8.66	10.12				
	NT2RP4001105	11.53	6.11	5.68	11.42	12.40	12.53	6.82	8.59	7.03				
	NT2RP4001110	4.14	2.11	2.03	3.53	3.73	5.22	7.74	9.16	5.7		*	+	
	NT2RP4001115	8.23	4.76	5.40	7.44	6.61	6.42	6.49	8.54	8.25				
	NT2RP4001117	5.86	2.61	3.66	4.84	5.68	5.67	6.82	7.82	11.35				
35	NT2RP4001122	4.53	2.89	4.44	5.52	5.25	6.14	3.92	5.33	5.22	*	+		
	NT2RP4001123	11.03	6.64	4.19	7.23	8.62	6.22	6.52	4.59	7.16				
	NT2RP4001126	12.30	8.14	5.35	14.50	10.35	10.40	6.7	7.95	9.08				
	NT2RP4001127	2.67	1.52	0.45	2.09	2.22	1.57	1.96	4.17	3				
	NT2RP4001138	3.41	2.11	1.63	1.48	2.64	1.74	2.14	3.24	4.17				
	NT2RP4001143	6.89	2.21	3.01	4.13	4.68	5.32	4.17	5.67	5.66				
40	NT2RP4001148	1.94	1.16	1.16	2.70	2.05	0.60	1.41	3.15	1.62				
	NT2RP4001149	4.34	2.11	2.80	3.19	3.00	3.41	3.12	4.58	4.05				
	NT2RP4001150	4.09	2.84	2.82	5.63	5.48	6.34	4.62	4.61	4.79	**	+	*	+
	NT2RP4001159	8.72	3.82	5.00	5.57	8.96	6.80	7.8	6.33	6.38				
	NT2RP4001162	3.97	2.49	1.88	3.46	2.36	3.14	3.98	2.29	2.75				
45	NT2RP4001170	9.81	5.75	5.29	2.68	3.96	2.23	2.4	2.44	1.3		*	-	
	NT2RP4001174	6.78	5.08	5.60	9.49	9.90	7.92	7.08	5.86	4.66	*	+		
	NT2RP4001175	19.07	9.74	10.40	16.34	17.86	15.79	8.78	8.58	11.27				
	NT2RP4001176	62.90	39.84	55.63	104.65	115.71	110.77	63.62	58.35	46.85	**	+		
	NT2RP4001184	10.39	5.65	5.39	5.95	4.48	5.41	4.76	4.78	4.24				
50	NT2RP4001198	10.79	4.11	5.82	13.69	9.03	11.21	14.64	14.06	13.84		*	+	
	NT2RP4001199	2.92	0.71	0.91	2.99	2.97	1.91	3.68	2.25	2.92				
	NT2RP4001206	13.96	4.32	7.41	11.41	10.25	10.46	8.73	9.26	10.42				
	NT2RP4001207	3.37	2.92	1.08	2.45	1.58	1.84	2.26	2.66	0.61				
	NT2RP4001210	2.36	1.47	2.10	3.13	2.39	1.71	1.5	2.49	2.3				
	NT2RP4001213	10.44	5.34	6.49	11.64	9.13	13.58	7.15	5.01	5.42				
55	NT2RP4001214	0.95	1.06	0.59	2.80	1.54	8.36	1.71	2.54	1.49		*	+	
	NT2RP4001219	2.55	2.66	2.86	4.42	15.66	4.45	5.58	7.03	5.57		**	+	

Table 286

5	NT2RP4000728	21.11	12.54	13.41	26.39	33.93	29.91	18.2	20.00	17.52	*	+		
	NT2RP4000737	2.29	1.59	0.36	2.95	3.74	3.56	1.99	4.28	1.59	*	+		
	NT2RP4000739	3.68	1.68	1.40	3.64	3.60	3.19	3.01	1.32	2.06				
	NT2RP4000749	4.61	2.23	2.17	5.43	5.08	3.32	3.77	2.84	2.99				
	NT2RP4000769	4.46	2.77	1.61	5.35	5.75	3.06	3.69	3.92	2.49				
	NT2RP4000774	7.04	3.62	4.69	6.48	7.03	5.14	4.99	3.77	3.67				
10	NT2RP4000781	1.78	1.82	2.45	2.48	1.82	2.08	1.95	1.67	1.08				
	NT2RP4000783	5.52	3.48	3.60	5.32	4.17	5.29	1.54	2.21	1.91		*	-	
	NT2RP4000787	(0.08)	0.27	0.06	0.45	0.09	1.07	0.1	0.13	-0.1				
	NT2RP4000788	7.00	4.42	3.89	7.56	7.52	5.50	5.26	4.25	3.66				
	NT2RP4000792	9.90	5.45	5.18	4.82	3.85	3.35	2.89	1.10	1.13		*	-	
15	NT2RP4000809	138.97	85.82	100.50	13.12	12.28	11.89	8.69	10.55	11.51	**	-	**	-
	NT2RP4000817	6.53	3.13	3.81	7.81	8.21	7.10	5.75	6.24	6.14	*	+		
	NT2RP4000821	10.40	5.88	5.97	8.60	9.00	10.24	19.32	14.83	13.61		*	+	
	NT2RP4000822	7.54	4.48	4.61	11.43	10.03	11.32	7.11	5.54	4.78	**	+		
	NT2RP4000823	6.10	4.87	14.52	6.50	4.58	4.69	17.58	17.55	14.17		**	+	
20	NT2RP4000831	4.53	2.70	1.65	4.00	4.27	4.75	3.68	4.83	3.77				
	NT2RP4000833	9.98	4.61	3.88	12.93	9.95	9.75	7.85	6.14	9.61				
	NT2RP4000837	16.84	7.67	8.19	4.27	7.04	6.55	7.9	6.72	7.63				
	NT2RP4000839	8.09	4.28	3.15	6.64	6.35	8.56	6.01	3.49	4.81				
	NT2RP4000846	7.97	4.70	3.74	7.70	5.83	5.14	6.12	4.09	4.55				
25	NT2RP4000848	5.78	2.64	3.11	8.90	6.26	8.65	7.07	7.56	8.46	*	+	*	+
	NT2RP4000855	3.22	3.08	1.54	2.41	2.92	2.82	2.82	2.57	2				
	NT2RP4000863	3.79	2.50	2.36	1.24	1.67	1.78	2	2.70	1.71				
	NT2RP4000865	9.55	7.40	5.94	26.23	26.54	18.52	8.98	8.90	8.56	**	+		
	NT2RP4000873	8.88	4.73	4.97	9.82	9.15	8.69	10.43	4.81	6.51				
	NT2RP4000874	5.60	3.25	3.18	4.02	6.09	6.60	5.15	3.17	5.54				
30	NT2RP4000875	10.06	7.69	6.92	10.24	9.60	8.28	5.61	5.34	4.98		*	-	
	NT2RP4000878	15.02	8.48	6.31	16.61	14.17	15.37	18.42	13.92	17				
	NT2RP4000879	1.68	0.79	0.77	1.38	2.21	2.54	2.35	2.03	1.86		*	+	
	NT2RP4000880	5.88	4.11	3.04	9.39	7.05	7.35	6.97	5.69	5.31	*	+		
35	NT2RP4000891	102.85	62.84	77.22	114.50	151.60	104.23	43.98	42.97	34.75		*	-	
	NT2RP4000894	8.78	5.12	4.69	6.91	6.62	9.49	7.97	4.83	7.88				
	NT2RP4000898	0.75	1.23	0.33	0.94	1.28	0.69	1.75	1.00	0.58				
	NT2RP4000899	14.91	8.73	9.27	8.87	7.17	6.06	2.92	6.91	6.96				
	NT2RP4000907	7.23	4.77	4.04	8.01	14.43	8.65	11.43	9.68	10.25		**	+	
	NT2RP4000908	3.70	3.82	2.81	5.39	5.05	5.27	4.11	5.22	3.41	**	+		
40	NT2RP4000910	11.95	5.36	6.97	10.03	8.98	9.73	9.64	9.49	7.69				
	NT2RP4000918	10.45	8.95	8.11	12.80	9.01	11.75	7.94	8.71	6.88				
	NT2RP4000925	1.77	2.18	1.68	2.08	2.56	3.09	1.91	2.37	0.93				
	NT2RP4000927	2.00	0.98	0.64	1.21	1.11	1.91	1.67	2.03	0.45				
	NT2RP4000928	8.63	5.13	3.60	5.86	6.72	6.51	5.18	4.85	6.75				
45	NT2RP4000929	1.61	1.10	1.06	1.59	2.36	1.14	0.96	1.23	1.92				
	NT2RP4000946	3.91	2.24	2.26	7.89	6.10	6.89	5.7	5.35	4.43	**	+	*	+
	NT2RP4000947	1.12	1.54	1.05	1.80	1.82	0.62	1.3	1.55	0.89				
	NT2RP4000949	16.12	8.67	10.24	5.88	3.51	5.79	19.02	19.45	15.95				
	NT2RP4000953	9.21	5.55	4.76	5.43	4.34	5.39	4.04	4.48	4.02				
	NT2RP4000959	16.07	16.16	17.01	17.30	15.74	18.65	13.76	14.61	12.03		*	-	
50	NT2RP4000962	4.28	2.72	4.02	3.76	4.20	2.99	2.02	3.10	1.89				
	NT2RP4000973	6.76	3.78	2.61	4.40	5.08	4.18	8.32	7.27	7.89				
	NT2RP4000975	4.74	2.41	1.77	5.26	4.90	3.72	2.88	4.71	4.07				
	NT2RP4000979	6.80	3.38	3.74	6.77	5.99	3.62	6.11	4.01	4.79				
	NT2RP4000984	3.24	3.46	2.61	2.85	2.49	5.25	1.35	3.81	1.22				
55	NT2RP4000986	3.13	2.19	3.27	2.70	3.05	3.24	3.2	4.03	2.69				
	NT2RP4000988	4.24	3.53	3.97	6.52	7.14	6.40	4.03	5.72	2.89	**	+		
	NT2RP4000989	4.55	3.53	3.49	5.18	3.51	4.95	4.91	5.46	4.69		*	+	

Table 285

	NT2RP4000453	7.28	6.16	3.48	2.35	2.43	4.15	1.8	4.72	0.91				
5	NT2RP4000455	1.01	1.01	1.48	2.29	2.70	1.92	2.22	2.27	0.83	*	+		
	NT2RP4000456	13.97	7.10	6.36	13.16	13.46	10.68	8.85	8.11	5.28				
	NT2RP4000457	6.68	4.82	2.84	3.69	4.73	3.69	4.6	3.98	5.62				
	NT2RP4000461	5.28	3.96	3.32	7.87	8.68	6.42	5.85	6.52	5.36	*	+		
	NT2RP4000462	8.07	4.05	4.23	7.49	8.39	11.75	6.93	5.29	4.06				
10	NT2RP4000463	9.18	6.18	6.85	10.59	9.85	9.05	5.78	4.84	4.27				
	NT2RP4000471	3.55	1.94	1.96	3.21	3.41	4.25	4.22	4.59	2.95				
	NT2RP4000472	3.05	2.42	1.96	12.20	8.76	6.84	4.36	5.24	4.11	*	+	*	+
	NT2RP4000476	1.50	1.02	0.85	12.49	11.85	10.88	21.84	18.65	17.71	**	+	**	+
	NT2RP4000480	15.36	6.51	5.30	5.47	9.87	5.81	7.44	7.54	5.87				
15	NT2RP4000481	3.47	2.35	0.78	2.35	2.92	2.36	3.06	3.89	4.07				
	NT2RP4000483	2.86	2.52	1.45	2.10	2.49	1.39	3.11	4.18	2.64				
	NT2RP4000487	3.11	1.79	1.56	6.59	4.70	2.73	3.7	3.87	2.46				
	NT2RP4000496	0.65	2.01	0.43	0.74	1.20	0.89	1.64	1.30	1.26				
	NT2RP4000497	6.68	4.62	5.43	14.85	10.68	12.20	7.76	11.46	5.67	**	+		
20	NT2RP4000498	4.09	1.89	2.15	3.59	3.39	3.97	3.69	5.45	2.91				
	NT2RP4000500	3.65	2.95	1.78	3.44	3.70	2.25	3.4	3.63	2.11				
	NT2RP4000507	15.14	8.22	5.69	11.50	10.49	7.06	7.7	7.22	9.04				
	NT2RP4000515	15.49	10.59	8.57	12.80	13.50	16.10	12.82	10.19	8.69				
	NT2RP4000516	7.24	4.39	3.65	20.66	19.29	17.91	10.11	9.21	8.83	**	+	*	+
25	NT2RP4000517	3.07	2.43	1.84	4.04	5.74	5.81	3.42	4.89	3.38	*	+		
	NT2RP4000518	4.18	1.91	2.39	4.28	2.50	2.78	3.19	3.42	2.91				
	NT2RP4000519	1.25	1.47	1.18	2.14	1.80	1.86	1.53	2.34	1.09	**	+		
	NT2RP4000524	0.66	1.08	0.33	1.66	1.94	1.79	1.87	1.81	1.62	**	+	**	+
	NT2RP4000528	1.96	2.16	0.43	1.52	2.71	2.98	1.9	3.84	1.18				
30	NT2RP4000537	40.32	18.87	17.18	18.72	15.16	10.99	14.21	10.18	11.8				
	NT2RP4000541	6.42	4.52	3.64	6.16	5.27	3.57	5.96	5.32	5.79				
	NT2RP4000543	7.15	4.38	3.94	5.71	5.28	6.49	7.13	6.85	7.19				
	NT2RP4000545	22.00	12.60	11.90	35.02	30.28	28.43	15.85	15.53	13.71	*	+		
	NT2RP4000546	3.49	2.74	2.72	5.16	6.84	5.20	2.65	5.26	4.13	*	+		
	NT2RP4000549	10.31	6.26	6.97	10.02	6.99	7.06	17.04	10.70	13.71				
35	NT2RP4000556	4.79	2.38	2.09	2.96	4.95	3.16	3.01	3.93	2.39				
	NT2RP4000557	2.43	1.89	1.59	3.06	2.06	2.13	1.6	1.76	2.34				
	NT2RP4000558	7.85	4.61	3.47	5.80	4.60	4.48	8.11	4.97	5.07				
	NT2RP4000560	11.62	8.43	5.62	16.38	11.32	8.62	10.3	8.86	6.76				
	NT2RP4000568	0.86	1.06	0.72	1.99	2.89	2.56	1.2	1.79	1.98	**	+	*	+
40	NT2RP4000583	9.91	5.21	4.91	9.30	13.09	14.53	6.79	5.52	7.23				
	NT2RP4000585	3.74	2.64	3.88	4.44	2.94	3.43	2.78	2.68	3.99				
	NT2RP4000588	1.78	1.61	0.91	2.23	3.68	2.01	2.78	3.01	2.89		**	+	
	NT2RP4000598	7.09	4.23	3.81	4.80	5.51	5.49	5.51	5.97	3.62				
	NT2RP4000599	1.53	1.26	0.87	1.24	1.41	1.06	0.44	2.70	0.51				
45	NT2RP4000603	11.90	6.03	3.85	6.61	6.16	3.84	4.98	5.10	6.79				
	NT2RP4000607	9.25	5.54	5.52	6.95	7.07	10.29	4.24	5.47	7.66				
	NT2RP4000614	18.95	12.78	10.17	25.67	26.47	23.13	9.33	11.19	9.77	*	+		
	NT2RP4000634	4.83	2.61	1.81	7.54	6.71	5.97	5.4	7.61	4.39	*	+		
	NT2RP4000638	3.55	2.37	1.27	3.88	3.82	3.28	2.34	4.08	2.48				
	NT2RP4000648	3.49	3.15	1.64	4.18	4.00	1.87	2.79	3.50	2.8				
50	NT2RP4000657	7.42	4.66	4.89	3.76	5.89	4.90	4.79	4.73	4.39				
	NT2RP4000691	3.57	4.48	4.25	6.09	7.82	5.58	5.65	7.17	5.49	*	+	*	+
	NT2RP4000697	11.06	7.17	4.24	7.59	7.47	5.97	4.38	5.33	7.55				
	NT2RP4000704	9.94	4.45	4.08	7.72	7.80	6.93	8.9	11.64	11.09				
	NT2RP4000710	39.78	22.43	20.25	37.57	42.17	34.47	22.39	29.16	28.71				
55	NT2RP4000713	3.09	1.40	0.88	3.21	4.08	3.16	3.3	5.18	2.97				
	NT2RP4000724	3.53	1.86	1.77	4.48	4.24	3.42	3.25	6.43	3.91				
	NT2RP4000725	4.59	2.50	2.14	3.16	3.33	2.21	3.39	4.06	2.51				

Table 284

	NT2RP4000175	26.66	17.23	19.20	12.23	15.62	11.17	16.22	18.62	19.97				
5	NT2RP4000180	17.71	15.54	16.60	7.75	7.76	10.71	9.21	10.11	9.68	**	-	**	-
	NT2RP4000185	14.57	9.35	-5.99	12.31	15.65	9.34	8.25	9.02	7.47				
	NT2RP4000192	9.26	5.09	4.80	6.32	4.48	3.65	4.83	4.74	4.23				
	NT2RP4000194	3.63	2.75	1.83	3.79	5.80	2.67	3.51	4.32	4.95				
	NT2RP4000196	8.18	4.81	3.10	7.96	7.13	5.03	5.27	5.97	5.49				
10	NT2RP4000210	28.53	18.46	17.26	28.89	37.05	27.38	24.22	22.19	25.95				
	NT2RP4000212	12.06	7.92	6.39	16.76	20.50	16.60	12.59	12.83	12.92	*	+		
	NT2RP4000214	10.71	7.74	6.94	13.03	16.29	15.56	10.1	11.28	8.66	*	+		
	NT2RP4000216	5.44	4.53	4.98	6.46	9.49	6.90	5.75	6.76	4.95				
	NT2RP4000218	7.33	2.22	2.67	4.98	4.33	3.81	3.81	4.58	6.27				
15	NT2RP4000223	19.92	13.17	10.28	22.13	21.62	13.05	22.62	26.76	25.86		*	+	
	NT2RP4000243	13.18	9.89	7.93	15.15	23.34	10.85	12.84	16.56	15.03				
	NT2RP4000246	33.96	22.95	19.51	28.17	27.99	24.14	21.88	39.67	28.61				
	NT2RP4000250	7.99	6.43	5.04	12.08	14.24	11.05	12.85	25.59	17.99	**	+	*	+
	NT2RP4000256	2.39	2.62	4.51	3.73	3.59	2.62	3.4	5.63	3.02				
20	NT2RP4000257	47.78	28.06	32.52	17.19	17.58	12.15	20.3	21.14	18.74	*	-		
	NT2RP4000259	4.57	3.53	4.63	12.50	13.85	8.56	9.95	10.96	10.32	*	+	**	+
	NT2RP4000261	4.69	3.90	2.69	4.69	4.12	2.59	6.07	3.27	3.23				
	NT2RP4000262	8.40	4.25	5.05	10.81	7.69	5.18	7.05	4.76	3.22				
	NT2RP4000263	2.39	2.26	1.46	3.24	1.78	2.52	2.31	2.43	1.67				
25	NT2RP4000280	19.84	10.94	16.02	14.51	20.53	17.86	16.38	15.79	14.33				
	NT2RP4000286	14.05	12.14	5.20	8.66	7.23	8.18	6.73	10.62	6.93				
	NT2RP4000290	4.20	3.07	2.79	5.43	3.58	4.59	3.38	3.10	2.4				
	NT2RP4000291	18.51	15.32	18.47	45.30	38.54	34.77	17.5	19.25	13.11	**	+		
	NT2RP4000301	2.59	1.81	1.04	2.23	2.98	3.54	2.54	3.49	1.63				
30	NT2RP4000312	4.56	1.79	4.33	4.54	4.75	3.56	5.14	2.41	5.06				
	NT2RP4000321	13.60	6.74	4.54	13.92	11.99	10.85	8.51	8.80	9.62				
	NT2RP4000323	3.58	2.53	1.59	2.86	3.50	3.23	2.71	3.60	1.23				
	NT2RP4000324	7.25	5.08	2.70	5.19	6.35	3.74	5.48	4.98	4				
	NT2RP4000334	13.97	11.43	12.75	30.03	27.15	21.64	10.28	10.30	9.71	**	+	*	-
	NT2RP4000343	4.98	3.25	2.65	4.86	5.56	3.68	3.76	4.39	3.15				
35	NT2RP4000348	3.02	1.79	1.77	4.45	3.35	4.09	4.17	3.46	2.74	*	+		
	NT2RP4000349	2.02	3.31	1.01	2.05	0.64	3.58	0.41	1.43	0.27				
	NT2RP4000355	10.07	4.28	4.14	7.89	8.66	7.17	5.76	4.78	6.28				
	NT2RP4000356	10.81	5.71	5.12	9.75	8.69	6.70	12.73	12.78	15.8		*	+	
	NT2RP4000360	5.76	3.41	2.25	11.67	15.48	9.10	8.87	7.21	7.44	*	+	*	+
40	NT2RP4000367	2.23	2.01	1.13	1.88	2.90	1.83	2.17	1.67	2.44				
	NT2RP4000370	4.54	3.75	1.61	3.50	4.39	3.20	3.15	3.31	3.03				
	NT2RP4000373	4.40	4.53	4.20	4.85	4.38	4.02	3.74	3.46	2.82		*	-	
	NT2RP4000376	3.46	3.35	3.32	5.35	3.36	3.31	2.76	4.60	2.39				
	NT2RP4000381	3.20	2.91	2.81	7.76	5.97	5.48	3.69	3.62	2.58	**	+		
45	NT2RP4000388	507.68	363.39	334.24	288.84	217.90	196.35	431.3	437.24	362.7				
	NT2RP4000390	19.01	14.68	11.68	24.99	29.51	23.19	15.68	13.59	14.64	*	+		
	NT2RP4000393	3.40	2.87	1.85	2.59	3.15	3.33	5.06	3.98	3.29				
	NT2RP4000398	5.34	4.23	2.50	10.36	14.48	10.01	6.8	5.94	5.69	*	+		
	NT2RP4000406	9.30	5.25	6.26	5.59	5.04	6.35	7.54	6.32	4.52				
	NT2RP4000407	5.98	4.41	3.78	8.29	7.16	4.70	4.32	5.68	5.13				
50	NT2RP4000413	1.40	1.18	0.62	0.72	1.57	3.58	1.37	2.49	1.36				
	NT2RP4000415	10.74	4.75	5.55	8.27	6.74	8.60	4.84	5.48	2.05				
	NT2RP4000417	7.49	5.67	3.62	5.24	6.05	4.58	5.78	5.18	6.52				
	NT2RP4000423	10.91	8.43	6.08	17.00	12.75	12.74	5.48	6.12	5.86	*	+		
	NT2RP4000424	4.48	2.86	1.81	7.46	7.77	6.37	5.69	7.35	4.76	**	+		
55	NT2RP4000447	13.10	8.03	11.15	9.03	13.44	9.03	6.38	5.33	5.62			*	-
	NT2RP4000448	2.34	1.79	0.84	4.19	6.84	6.98	5.24	4.20	3.76	*	+	*	+
	NT2RP4000449	2.70	2.01	2.13	2.07	1.89	2.22	2.41	2.65	1.44				

Table 283

	NT2RP3004557	9.04	5.56	6.56	5.65	4.56	3.38	5.82	5.13	3.59			
	NT2RP3004561	5.68	3.44	3.35	5.27	5.92	3.88	4.61	5.03	4.06			
5	NT2RP3004566	6.63	6.29	6.33	12.53	11.01	9.47	7.43	8.46	13.57	**	+	
	NT2RP3004569	6.44	5.29	4.60	10.37	11.99	10.11	4.46	4.55	4.39	**	+	
	NT2RP3004572	3.83	3.21	2.73	4.62	5.78	5.28	4.26	4.30	2.97	*	+	
	NT2RP3004578	5.21	3.44	2.27	5.01	7.11	5.48	3.71	3.96	4.42			
	NT2RP3004584	3.59	3.64	3.56	3.31	4.74	4.86	3.85	3.43	4.22			
10	NT2RP3004588	3.87	2.70	2.67	8.15	6.21	6.68	4.64	5.48	4.37	**	+	* +
	NT2RP3004594	7.86	6.82	6.37	5.22	4.81	5.30	4.15	4.02	2.13	*	-	* -
	NT2RP3004603	60.30	35.19	34.71	45.07	50.01	29.71	17.9	21.98	18.08		*	-
	NT2RP3004612	6.20	3.05	3.45	4.40	4.92	2.76	4.05	3.39	3.11			
	NT2RP3004617	3.07	2.70	1.70	1.60	2.01	3.22	2.53	2.44	1.96			
15	NT2RP3004618	3.95	2.90	2.07	5.51	5.52	3.64	3.14	3.14	4.18			
	NT2RP3004625	5.48	4.10	2.95	5.75	7.50	5.56	7.41	6.90	5.44			
	NT2RP3004635	4.31	4.50	4.46	4.30	6.48	5.74	5.58	3.86	3.99			
	NT2RP3004640	3.88	3.08	3.28	7.49	7.45	6.73	5.96	5.47	4.27	**	+	* +
	NT2RP3004642	10.28	8.51	8.84	14.09	13.53	15.70	10	10.58	5.55	**	+	
20	NT2RP3004647	7.16	4.79	5.37	9.93	6.54	8.91	7.81	5.99	5.6			
	NT2RP3004652	9.07	6.60	3.76	13.15	12.30	9.92	7.24	7.33	3.44	*	+	
	NT2RP3004669	8.16	5.80	4.33	5.00	7.93	5.74	5.7	5.73	5.33			
	NT2RP3004670	14.41	12.39	9.32	16.29	20.04	15.04	13.36	13.59	15.01			
	NT2RP4000008	15.39	10.91	11.09	13.50	10.87	9.28	9.4	8.75	8.85			
25	NT2RP4000018	9.99	5.44	8.54	9.01	5.02	7.90	7.84	6.47	7.74			
	NT2RP4000023	5.20	4.00	3.38	3.86	2.64	2.61	3.51	4.32	2.67			
	NT2RP4000025	5.36	5.89	4.96	8.91	15.04	11.95	12.96	16.75	13.7	*	+	** +
	NT2RP4000035	8.26	5.47	5.42	13.88	11.54	12.72	5.97	11.43	5.65	**	+	
	NT2RP4000041	8.69	5.46	1.79	1.69	4.25	2.76	4.28	5.58	4.93			
30	NT2RP4000049	4.05	2.09	2.36	3.68	4.19	3.53	5.9	5.73	3.33			
	NT2RP4000050	3.62	2.75	1.71	2.29	3.50	3.25	3.01	5.38	3.14			
	NT2RP4000051	7.84	3.90	4.64	5.71	7.58	5.48	5.27	7.15	5.15			
	NT2RP4000063	4.66	2.43	2.44	3.26	2.94	4.77	3.68	5.96	2.61			
	NT2RP4000065	4.21	2.76	2.69	4.09	3.65	3.77	3.32	3.08	2.24			
35	NT2RP4000070	3.16	2.60	2.02	6.63	8.48	9.49	3.2	4.92	3.34	**	+	
	NT2RP4000074	1.25	0.65	0.45	1.09	0.95	1.43	1.92	3.35	1.24			
	NT2RP4000078	19.45	8.95	8.65	15.20	11.49	10.74	9.98	6.63	6.98			
	NT2RP4000080	16.31	10.55	9.31	16.83	24.18	15.57	14.36	10.43	16.69			
	NT2RP4000099	48.25	34.08	34.96	222.14	203.11	165.35	108.2	86.72	64.03	**	+	* +
40	NT2RP4000102	1.59	3.03	0.75	2.02	3.06	3.50	2.33	2.26	2.57			
	NT2RP4000103	2.96	1.87	1.69	2.51	4.74	2.46	2.75	4.73	2.41			
	NT2RP4000108	7.32	4.36	4.82	47.03	44.25	37.96	49.26	38.51	49.37	**	+	** +
	NT2RP4000109	12.97	8.34	8.98	9.50	12.20	12.85	13.79	10.89	9.27			
	NT2RP4000111	1.66	4.14	1.76	3.30	2.22	1.71	2.22	1.42	3.11			
	NT2RP4000112	12.62	5.96	5.20	13.14	12.78	6.27	9.14	9.28	9.82			
45	NT2RP4000115	6.69	4.45	3.10	4.28	5.71	3.35	6.12	5.23	4.95			
	NT2RP4000129	5.85	2.83	2.30	2.80	3.92	3.49	3.8	3.85	2.88			
	NT2RP4000137	6.85	6.38	5.53	4.82	7.68	8.16	4.3	6.03	5.81			
	NT2RP4000138	31.16	22.51	24.42	13.11	12.17	10.03	14.81	14.41	15.27	**	-	* -
	NT2RP4000141	4.89	2.65	2.93	4.06	3.52	4.29	2.76	4.18	2.03			
50	NT2RP4000147	2.17	1.29	1.74	2.55	2.46	3.03	2.68	3.29	2.54	*	+	* +
	NT2RP4000150	7.08	4.20	5.06	8.60	7.56	6.25	7.64	8.70	6.48			
	NT2RP4000151	7.65	4.77	3.15	5.40	5.42	4.70	5.71	4.77	7.3			
	NT2RP4000157	47.42	28.18	24.63	140.24	151.70	90.24	64.55	61.24	48.04	**	+	* +
	NT2RP4000159	2.50	1.76	1.15	1.15	1.62	2.34	1.61	2.61	1.83			
55	NT2RP4000163	26.39	20.86	16.59	7.91	9.36	8.09	5.61	5.24	4.41	*	-	** -
	NT2RP4000167	3.26	3.04	2.67	3.80	3.99	4.24	2.64	3.85	3.17	**	+	
	NT2RP4000171	7.53	5.74	5.41	5.89	7.46	4.62	5.54	5.19	6.82			

Table 282

5	NT2RP3004262	4.35	2.96	2.85	2.71	3.57	4.45	4.01	4.72	3.41				
	NT2RP3004275	3.72	3.04	2.37	3.29	3.02	3.38	3.39	4.75	1.04				
	NT2RP3004282	12.87	5.01	5.72	9.16	11.91	6.32	7.38	7.58	6.69				
	NT2RP3004289	3.01	2.85	1.46	6.88	5.77	3.72	2.35	3.31	3.68	*	+		
	NT2RP3004294	7.18	3.41	2.73	24.46	29.15	28.18	20.58	15.67	20.34	**	+	**	+
	NT2RP3004298	7.07	5.08	3.77	5.00	5.97	6.16	6.4	6.06	5.61				
10	NT2RP3004309	10.96	7.28	6.61	7.01	8.68	7.42	5.52	6.85	6.57				
	NT2RP3004321	11.18	6.12	7.27	9.56	8.71	10.32	7.19	8.23	10.39				
	NT2RP3004322	3.28	2.42	1.89	3.12	2.58	3.70	3.77	3.09	3.39				
	NT2RP3004332	6.32	6.72	6.36	11.24	8.54	10.03	4.86	8.82	5.48	*	+		
	NT2RP3004334	4.49	2.34	2.27	5.43	4.10	3.66	2.44	1.92	2.32				
15	NT2RP3004336	5.86	3.72	2.08	6.83	9.08	6.19	5.13	6.87	5.49				
	NT2RP3004338	11.56	5.52	9.71	8.36	5.67	6.93	5.31	4.61	6.32				
	NT2RP3004341	2.24	1.74	1.67	2.56	2.48	3.60	1.13	2.35	3.45				
	NT2RP3004345	3.27	3.23	2.25	3.71	4.02	3.88	3.2	3.07	4.38				
	NT2RP3004348	8.53	5.32	16.83	14.49	13.97	11.82	7.76	7.80	9.23	**	+		
20	NT2RP3004349	10.22	7.24	8.20	12.70	11.94	13.01	6.98	7.06	5.47	*	+		
	NT2RP3004355	6.08	5.70	3.65	5.80	6.46	7.00	4.88	5.01	4.97				
	NT2RP3004356	13.62	7.29	6.71	12.35	15.04	10.32	9.71	9.44	9.13				
	NT2RP3004360	7.52	3.61	3.49	4.81	4.04	4.08	2.07	3.17	4.82				
	NT2RP3004361	16.01	7.31	5.66	15.99	14.58	14.13	4.38	5.01	4.13				
25	NT2RP3004374	7.91	4.13	3.84	7.91	7.91	7.64	5.99	5.39	5.89				
	NT2RP3004378	26.21	17.19	14.59	10.81	12.69	11.18	6.13	10.86	9.07		*		
	NT2RP3004399	2.04	2.65	1.39	1.42	2.99	2.67	1.58	2.38	2.75				
	NT2RP3004405	3.95	3.77	2.00	4.65	7.05	3.79	3.22	5.96	4.47				
	NT2RP3004406	7.20	4.61	5.55	5.61	8.40	5.80	5.82	7.89	6.47				
	NT2RP3004411	7.77	3.85	3.09	16.41	12.18	7.61	7.04	7.47	10.13				
30	NT2RP3004424	4.60	1.42	1.67	3.96	3.79	2.00	1.27	3.09	4.78				
	NT2RP3004428	7.15	4.01	3.24	6.42	5.85	3.58	6.97	6.90	7.98				
	NT2RP3004432	3.82	2.57	0.97	7.56	9.25	7.81	7.72	10.80	9.98	**	+	**	+
	NT2RP3004434	9.49	5.09	3.75	6.31	8.59	6.98	5.23	4.83	5.64				
	NT2RP3004446	6.23	5.35	3.39	6.60	5.96	4.57	2.58	4.37	4.71				
35	NT2RP3004451	3.49	1.02	1.26	4.55	6.79	4.04	2.13	3.69	4.46				
	NT2RP3004454	3.00	1.25	1.36	2.36	2.23	1.93	1.66	2.42	2.5				
	NT2RP3004466	16.12	6.82	7.66	12.66	11.01	12.35	11.52	8.75	10.08				
	NT2RP3004470	8.70	6.35	3.18	11.68	12.19	10.86	7.44	7.38	5.56	*	+		
	NT2RP3004472	1.89	2.60	1.02	4.08	3.19	3.82	2.45	1.91	1.78	*	+		
40	NT2RP3004475	4.99	3.80	4.98	4.54	5.61	3.71	4.55	5.07	4.35				
	NT2RP3004480	7.66	5.39	3.59	15.02	14.38	12.51	8.01	7.48	6.29	**	+		
	NT2RP3004481	4.24	6.01	3.44	3.84	4.84	6.10	5.51	4.88	3.41				
	NT2RP3004490	1.09	1.00	1.30	1.59	2.17	1.90	1.13	0.94	0.16	*	+		
	NT2RP3004496	11.99	5.64	6.80	14.82	15.35	7.87	12.41	15.48	10.73				
45	NT2RP3004498	10.57	6.90	5.91	5.39	8.13	7.76	7.22	4.55	5.58				
	NT2RP3004503	8.32	5.77	4.24	17.06	17.79	15.82	8.93	7.92	6.72	**	+		
	NT2RP3004504	16.66	9.32	8.13	4.90	5.37	6.99	5.11	6.36	4.24				
	NT2RP3004505	8.72	5.28	4.61	4.26	5.67	7.97	8.11	8.94	7.62				
	NT2RP3004507	4.86	3.25	3.44	5.31	4.59	4.43	2.27	3.02	3.06				
	NT2RP3004519	3.79	1.12	1.28	2.61	2.20	3.15	1.55	1.93	1.88				
50	NT2RP3004524	1.80	1.60	2.36	2.58	1.67	4.26	2.22	2.30	1.3				
	NT2RP3004527	1.16	0.95	0.83	1.29	0.90	1.98	0.25	0.54	0.6		*		
	NT2RP3004534	5.79	3.52	3.93	3.26	4.89	7.19	3.48	2.92	3				
	NT2RP3004539	14.05	8.61	6.22	8.74	9.46	10.33	9.59	6.77	9.38				
	NT2RP3004541	4.42	3.07	2.91	2.08	4.14	2.58	3.82	4.01	3.83				
	NT2RP3004544	9.72	3.68	2.35	4.38	6.86	7.17	4.05	5.81	5.64				
55	NT2RP3004551	3.07	2.54	2.87	4.55	5.33	3.79	4.49	4.42	3.29	*	+	*	+
	NT2RP3004552	11.09	5.45	4.69	3.94	6.09	8.11	5.03	5.45	2.94				

Table 281

	NT2RP3003989	2.69	2.90	1.66	1.97	4.23	17.93	2.15	5.56	2.09				
5	NT2RP3003992	4.45	3.19	2.09	6.85	5.45	5.48	2.46	5.01	2.52	*	+		
	NT2RP3004000	2.21	2.96	1.05	1.76	3.78	2.06	4.87	2.93	3.16				
	NT2RP3004001	10.03	7.36	4.34	11.63	8.96	9.72	6.39	7.58	6.18				
	NT2RP3004005	2.84	1.39	1.85	4.23	3.15	3.89	6.12	4.26	2	*	+		
	NT2RP3004013	12.35	8.49	6.06	13.19	14.63	10.33	6.81	8.18	5.23				
10	NT2RP3004016	4.50	2.25	1.85	4.36	3.71	4.81	2.81	2.48	3.43				
	NT2RP3004025	4.30	3.53	3.53	4.99	6.65	6.46	4.38	6.03	4.27	*	+		
	NT2RP3004030	22.90	14.65	17.74	29.69	32.04	29.24	22.51	18.90	19.9	**	+		
	NT2RP3004041	2.52	1.89	2.73	9.78	7.34	7.80	4.71	4.38	4.76	**	+	**	+
	NT2RP3004042	14.33	10.61	5.39	8.88	10.41	10.70	11.54	9.64	11.73				
15	NT2RP3004044	21.83	11.12	9.61	8.22	9.50	8.35	6.17	5.06	6.39				
	NT2RP3004051	10.03	6.48	4.32	11.50	10.92	8.70	7.09	5.39	5.97				
	NT2RP3004052	8.89	3.73	4.41	8.80	8.69	8.41	6.86	4.66	5.92				
	NT2RP3004053	30.17	20.41	22.51	39.10	49.24	42.11	31.51	31.71	33.47	*	+		
	NT2RP3004055	4.37	1.71	1.44	3.41	6.47	4.74	2.67	3.05	2.47				
20	NT2RP3004059	4.35	3.84	2.26	4.57	5.40	6.36	4.38	3.95	3.58				
	NT2RP3004063	3.19	5.38	4.25	5.25	3.73	4.82	2.48	4.55	2.33				
	NT2RP3004067	20.37	6.61	6.47	9.24	9.55	7.82	8.89	7.62	7.01				
	NT2RP3004070	5.14	4.09	2.46	6.23	5.56	5.86	3.96	3.22	4.36				
	NT2RP3004075	4.89	3.98	3.09	4.61	4.46	5.82	3.77	3.33	3.83				
	NT2RP3004078	6.60	3.72	3.12	5.82	6.46	5.79	5.42	4.95	4.97				
25	NT2RP3004083	2.32	2.07	2.04	35.55	41.35	31.65	20.9	19.75	24.51	**	+	**	+
	NT2RP3004084	4.82	3.89	2.80	2.32	2.21	5.07	2.3	4.34	3.24				
	NT2RP3004087	6.30	4.80	3.92	7.31	7.31	7.55	5.02	5.55	6.07	*	+		
	NT2RP3004090	3.22	2.13	1.57	4.35	5.08	3.83	3.16	6.01	4.35	*	+		
	NT2RP3004093	5.89	4.55	3.16	7.72	8.34	6.85	6.58	5.64	6.63	*	+		
30	NT2RP3004095	14.57	8.24	7.88	13.27	13.82	13.04	10.11	8.74	11.47				
	NT2RP3004102	11.19	6.90	6.93	9.17	11.74	10.70	9.42	7.28	9.35				
	NT2RP3004110	34.95	22.41	23.25	26.04	28.26	24.02	16.77	18.06	22.74				
	NT2RP3004119	6.91	5.16	5.08	8.05	6.96	6.49	5.73	4.85	4.73				
	NT2RP3004125	14.03	10.35	8.98	14.12	16.80	14.86	13.91	11.06	10.62				
35	NT2RP3004129	3.44	1.56	2.05	2.41	2.99	3.58	2.35	2.48	1.77				
	NT2RP3004130	3.67	2.75	3.57	6.28	6.18	5.89	7.37	7.97	5.85	**	+	**	+
	NT2RP3004133	8.07	5.45	4.56	6.17	4.98	5.72	6.99	6.13	6.19				
	NT2RP3004145	6.56	4.08	2.26	3.88	4.54	4.28	2.91	4.84	3.57				
	NT2RP3004148	7.79	6.05	5.54	5.61	5.84	7.93	7.7	7.31	5.13				
	NT2RP3004155	3.99	4.60	2.60	5.64	5.29	6.17	3.4	3.66	2.7	*	+		
40	NT2RP3004165	9.52	6.71	6.33	12.69	13.98	12.98	6.82	6.51	5.79	**	+		
	NT2RP3004179	4.17	3.60	3.22	5.35	6.25	6.22	3.75	3.01	3.75	**	+		
	NT2RP3004185	2.33	0.68	1.31	1.91	1.20	2.96	1.8	2.34	1.86				
	NT2RP3004188	8.37	4.08	5.91	11.26	11.20	6.76	4.54	7.20	6.27				
	NT2RP3004189	14.04	5.66	6.06	7.02	12.29	6.24	4.85	4.58	5.6				
45	NT2RP3004190	11.54	5.42	6.63	7.75	12.77	11.72	5.49	4.47	5.81				
	NT2RP3004191	10.44	9.83	8.83	14.00	14.26	11.80	12.41	10.04	10.36	*	+		
	NT2RP3004202	2.35	2.27	2.03	3.51	4.57	3.29	3.6	3.97	5.67	*	+	*	+
	NT2RP3004205	10.83	6.54	6.41	8.47	10.58	6.84	7.02	6.54	6.67				
	NT2RP3004206	3.85	2.53	2.95	2.95	3.06	3.06	4.12	2.99	2.57				
	NT2RP3004207	4.93	2.79	3.03	4.73	4.14	4.86	4.28	4.10	5.09				
50	NT2RP3004209	4.91	2.40	2.89	6.87	6.50	4.96	4.96	5.23	4.63	*	+		
	NT2RP3004215	3.55	2.78	2.14	12.42	8.20	7.94	3.86	4.27	5.18	*	+	*	+
	NT2RP3004219	16.93	6.45	7.83	7.64	9.11	7.10	7.36	6.55	7.25				
	NT2RP3004242	5.13	4.26	3.60	4.45	5.10	4.52	4.84	3.47	2.95				
	NT2RP3004246	4.82	4.45	3.64	5.22	7.08	6.18	4.56	5.99	5.39	*	+		
55	NT2RP3004253	1.98	2.17	2.49	2.39	1.93	2.99	2.38	3.72	5.59				
	NT2RP3004258	11.77	7.63	9.50	10.32	13.55	13.92	4.51	6.56	5.46		*	-	

Table 280

	NT2RP3003729	3.69	2.88	2.55	4.06	4.92	3.98	2.8	3.60	3.35	*	+		
5	NT2RP3003731	6.61	4.33	5.75	7.10	14.90	8.06	5.99	7.15	5.75				
	NT2RP3003740	4.78	3.50	4.29	5.32	3.89	4.79	4.16	4.89	3.61				
	NT2RP3003746	5.36	3.49	2.71	5.20	7.52	3.17	3.94	3.31	4.02				
	NT2RP3003749	0.76	0.62	0.17	0.29	1.19	1.12	0.64	1.30	0.75				
	NT2RP3003754	5.00	3.26	5.25	7.46	7.69	6.19	5.46	4.91	4.55	*	+		
10	NT2RP3003759	1.70	0.69	0.73	1.39	1.06	0.48	0.73	2.09	2.41				
	NT2RP3003764	7.97	5.68	5.63	6.40	8.69	7.67	5.36	5.99	4.9				
	NT2RP3003766	4.56	2.73	2.99	3.97	4.19	3.87	3.96	3.75	3.32				
	NT2RP3003767	6.96	5.70	6.63	13.57	9.41	11.81	7.79	9.76	8.37	*	+	*	+
	NT2RP3003778	5.19	3.99	4.33	9.90	11.58	8.75	5.62	5.86	5.15	**	+		
15	NT2RP3003779	13.01	5.97	4.99	6.05	7.93	6.85	7.17	5.72	8.58				
	NT2RP3003783	19.26	10.08	8.20	11.73	11.20	13.62	12.33	9.52	7.82				
	NT2RP3003787	4.90	2.40	2.22	2.44	3.52	4.85	2.78	3.53	7.22				
	NT2RP3003789	5.36	4.73	2.56	3.44	7.01	5.23	5.4	5.55	4.62				
	NT2RP3003795	2.17	1.85	1.40	3.14	2.08	3.57	2.46	3.18	2.41				
20	NT2RP3003799	2.89	2.29	1.32	1.87	1.75	2.53	1.45	2.24	2.66				
	NT2RP3003800	3.51	2.88	4.22	3.79	5.81	4.55	3.66	3.45	2.49				
	NT2RP3003805	6.47	3.37	3.41	4.89	4.12	5.73	3.59	4.60	4.09				
	NT2RP3003809	5.03	1.78	2.92	4.79	3.39	3.28	1.85	3.89	3.58				
	NT2RP3003819	20.93	12.43	10.20	22.69	23.35	18.68	16.05	13.33	11.82				
	NT2RP3003824	12.10	8.20	9.56	14.53	12.56	14.16	10.06	10.73	7.38	*	+		
25	NT2RP3003825	22.51	14.11	14.65	13.44	18.74	15.00	10.89	9.86	10.89				
	NT2RP3003828	3.66	3.06	2.75	5.51	4.72	4.12	2.65	4.12	4.14	*	+		
	NT2RP3003831	2.13	2.74	2.94	4.32	4.71	5.94	3.1	4.50	4.33	*	+		
	NT2RP3003833	5.17	2.54	2.51	3.72	3.00	5.07	4.52	4.42	4				
	NT2RP3003836	7.43	5.49	5.12	9.64	6.79	8.16	7.54	6.97	9.43				
30	NT2RP3003842	17.19	8.40	7.68	16.76	16.34	13.12	12.09	8.43	8.61				
	NT2RP3003843	11.40	7.50	6.65	20.59	22.26	19.09	11.26	10.84	11.37	**	+		
	NT2RP3003844	12.70	8.55	6.42	7.70	6.74	8.49	13.96	12.46	12.2				
	NT2RP3003846	3.76	1.97	2.48	4.49	3.48	4.92	2.73	3.31	3.38				
	NT2RP3003849	4.75	3.02	2.95	4.08	4.65	4.41	2.89	4.41	5.12				
35	NT2RP3003862	8.19	5.27	4.97	5.73	7.14	6.59	9.21	6.75	9.43				
	NT2RP3003870	8.87	6.42	4.81	9.09	8.35	8.66	8.21	7.03	8.25				
	NT2RP3003874	4.83	4.91	4.32	6.66	5.96	5.92	4.88	5.78	3.78	**	+		
	NT2RP3003876	8.40	4.71	3.53	8.21	6.66	5.04	3.88	4.35	5.13				
	NT2RP3003880	3.42	3.11	2.28	6.01	6.99	4.51	4.71	5.26	4.07	*	+	*	+
	NT2RP3003889	1.46	1.88	0.92	1.03	3.20	2.06	0.85	2.31	2.72				
40	NT2RP3003891	1.54	2.30	0.87	1.75	2.99	2.00	1.08	2.80	2.25				
	NT2RP3003914	7.95	4.51	4.21	5.57	7.65	7.02	5.69	6.39	7.2				
	NT2RP3003915	1.86	2.20	1.19	1.63	2.60	2.36	2.19	3.03	2.1				
	NT2RP3003918	5.05	3.66	2.14	2.83	4.62	2.98	3.63	5.42	5.25				
45	NT2RP3003920	4.98	4.36	2.71	6.50	6.25	5.72	5.51	6.91	3.85	*	+		
	NT2RP3003924	6.49	3.55	2.01	7.69	8.02	5.14	4.31	3.95	7.6				
	NT2RP3003932	3.65	2.42	1.71	4.82	7.98	3.41	2.85	3.76	4.41				
	NT2RP3003939	2.69	1.67	1.95	3.86	3.92	3.18	2.41	3.31	2.98	*	+		
	NT2RP3003940	15.51	8.52	7.81	11.47	11.25	8.35	8.68	9.97	7.23				
	NT2RP3003943	3.63	3.38	2.60	2.90	3.77	1.83	2.48	3.48	4.35				
50	NT2RP3003959	2.34	2.12	1.61	3.04	4.84	3.82	2.42	3.16	4.93	*	+		
	NT2RP3003963	6.98	5.43	4.54	7.42	7.40	5.93	6.05	7.92	6.84				
	NT2RP3003965	44.37	24.77	31.74	35.84	34.50	26.36	12.7	12.05	15.03		*	-	
	NT2RP3003972	14.33	10.15	6.83	27.44	20.29	23.76	23.62	15.59	17.39	*	+		
	NT2RP3003973	8.15	5.02	3.70	7.18	5.27	4.94	5.3	5.61	3.97				
55	NT2RP3003979	11.32	8.28	4.38	9.43	15.88	13.30	10.9	7.26	6.82				
	NT2RP3003980	10.84	7.99	7.63	8.16	9.43	9.50	5.75	7.95	4.2				
	NT2RP3003982	1.33	3.01	1.15	1.21	2.15	2.04	1.21	3.30	0.58				

Table 279

	NT2RP3003462	4.12	2.91	3.40	4.80	6.31	3.84	3.87	5.08	5.3				
	NT2RP3003464	2.09	1.93	2.25	2.69	2.89	1.33	2.08	3.11	2.15				
5	NT2RP3003469	3.14	2.14	3.25	2.94	4.36	3.12	4.25	4.89	4.48		*	+	
	NT2RP3003473	89.05	73.31	85.12	31.82	60.74	51.48	32.07	37.27	32.95	*	-	**	-
	NT2RP3003474	3.72	1.64	1.41	2.81	4.68	2.60	1.76	1.83	5				
	NT2RP3003475	5.61	2.84	3.02	4.26	5.48	3.96	3.12	2.86	5.38				
10	NT2RP3003490	2.57	1.77	0.90	2.92	3.66	2.60	1.94	2.99	7.73				
	NT2RP3003491	3.82	1.31	1.56	3.52	3.19	3.71	1.08	3.23	2.23				
	NT2RP3003493	32.32	24.24	22.86	18.58	22.23	21.78	10.29	22.93	16.25				
	NT2RP3003500	1.40	1.72	1.09	3.53	3.58	2.03	2.25	3.61	2.95	*	+	*	+
	NT2RP3003527	2.93	1.02	1.39	2.26	3.40	1.33	1.5	4.37	4.13				
15	NT2RP3003532	6.83	4.04	4.22	15.20	17.07	14.08	8.23	6.65	7.28	**	+		
	NT2RP3003535	1.58	1.03	0.30	1.85	1.07	0.98	1.62	1.27	0.97				
	NT2RP3003536	2.90	2.77	1.64	5.15	3.92	4.74	3.97	3.71	2.81	*	+		
	NT2RP3003543	4.72	4.39	3.25	5.41	8.08	7.02	5.2	5.49	1.98	*	+		
	NT2RP3003549	2.71	2.81	2.37	2.41	3.79	4.08	3.3	2.30	1.66				
20	NT2RP3003552	1.05	1.06	0.00	1.19	1.29	1.21	0.42	0.50	0.79				
	NT2RP3003555	7.69	3.49	4.38	7.36	8.38	9.29	5.4	5.33	4.02				
	NT2RP3003559	2.48	1.02	1.13	2.42	2.15	3.37	1.46	1.90	0.56				
	NT2RP3003564	6.10	3.28	3.23	6.06	5.72	4.12	4.46	3.78	4.48				
	NT2RP3003572	4.33	3.51	2.66	3.48	4.50	3.26	3.67	4.32	2.39				
25	NT2RP3003576	14.59	6.63	6.37	16.23	21.96	19.84	10.82	10.96	8.97	*	+		
	NT2RP3003587	15.06	8.22	7.88	8.40	8.95	10.51	3.86	6.31	3.33				
	NT2RP3003589	14.90	11.19	8.98	10.98	18.16	17.00	16.77	16.70	14.61				
	NT2RP3003592	6.07	3.40	4.66	3.72	5.45	5.40	3.54	4.53	3.89				
	NT2RP3003593	5.28	1.75	2.13	3.76	3.86	7.00	3.06	4.00	2.78				
	NT2RP3003614	14.05	8.27	10.10	10.29	8.15	9.17	8.06	7.21	4.02				
30	NT2RP3003621	3.29	1.07	1.69	2.23	2.27	2.45	2.08	2.77	2.99				
	NT2RP3003625	11.53	5.52	5.48	9.50	9.71	7.13	7.18	5.14	5.56				
	NT2RP3003627	12.05	7.44	6.80	53.97	42.81	41.76	14.96	15.18	18.14	**	+	*	+
	NT2RP3003636	5.65	3.72	2.95	5.93	6.64	5.54	5.93	5.72	5.63				
	NT2RP3003642	10.88	8.03	6.37	13.82	13.96	17.20	12.37	12.40	16.41	*	+	*	+
35	NT2RP3003645	4.17	3.33	1.50	5.78	5.31	6.65	5.06	5.99	4.7	*	+		
	NT2RP3003648	3.24	3.31	3.16	4.15	4.43	3.91	5.07	3.21	3.18	**	+		
	NT2RP3003649	1.14	1.88	2.86	2.19	4.90	3.66	0.71	3.92	1.08				
	NT2RP3003650	8.11	4.45	2.20	4.63	4.76	4.42	3.39	3.56	3.36				
	NT2RP3003656	5.22	3.74	1.88	3.30	4.87	4.62	3.45	3.40	2.71				
	NT2RP3003659	7.45	4.72	4.52	4.36	6.73	4.25	3.17	3.21	3.03				
40	NT2RP3003662	9.17	7.44	5.08	10.50	15.08	10.64	9.44	8.85	8.35				
	NT2RP3003664	8.73	4.21	6.55	11.31	14.75	9.76	9.24	8.50	9.85				
	NT2RP3003665	1.46	2.31	3.07	2.00	3.22	2.55	1.63	2.93	1.01				
	NT2RP3003671	3.15	3.24	2.25	2.59	7.96	5.47	2.17	4.14	1.93				
	NT2RP3003672	4.15	3.09	2.96	4.72	7.37	5.47	2.79	4.69	3.17	*	+		
45	NT2RP3003673	4.51	3.32	1.35	5.41	6.14	2.58	4.36	4.67	3.13				
	NT2RP3003679	34.38	42.38	35.15	32.46	39.83	37.84	41.64	35.07	42.5				
	NT2RP3003680	6.95	3.40	1.56	4.84	3.86	4.38	2.61	3.70	3.96				
	NT2RP3003686	5.14	3.55	2.82	3.79	4.38	5.04	4.26	3.62	2.84				
	NT2RP3003689	3.80	2.46	2.57	6.17	7.73	5.84	3.57	4.94	3.47	**	+		
50	NT2RP3003697	1.90	2.24	1.34	1.76	2.19	2.72	2.08	3.11	1.51				
	NT2RP3003701	1.92	1.12	1.36	1.56	1.36	1.59	2.02	2.99	1.34				
	NT2RP3003704	5.17	3.39	3.77	6.61	6.98	7.53	4.92	5.10	3.69	**	+		
	NT2RP3003714	3.30	1.91	1.74	4.60	3.93	3.09	3.44	3.54	1.64				
	NT2RP3003716	2.44	2.40	1.34	4.13	2.42	3.98	2.31	2.88	2.92				
	NT2RP3003721	4.90	3.12	2.28	4.84	6.16	4.98	4.29	3.50	4.65				
55	NT2RP3003722	8.02	5.81	5.39	6.08	4.24	4.20	2.09	3.67	2.7		*	-	
	NT2RP3003726	6.59	6.25	3.44	4.38	3.30	5.00	5.53	4.21	4.73				

Table 278

5	NT2RP3003139	2.43	1.97	1.82	4.72	6.45	3.81	3.26	3.26	4.15	*	+	*	+
	NT2RP3003145	2.66	3.16	2.32	3.58	4.86	4.52	5.45	3.67	3.72	*	+		
	NT2RP3003150	4.45	3.91	-3.35	3.70	3.28	5.66	5.36	4.59	2.96				
	NT2RP3003157	15.45	8.45	11.15	23.44	27.58	18.86	11.74	13.90	10.21	*	+		
	NT2RP3003185	3.41	2.15	1.16	2.42	3.21	3.33	3.63	2.51	4.07				
	NT2RP3003193	5.13	4.24	4.83	11.32	20.09	13.42	6.1	6.95	8.42	*	+	*	+
10	NT2RP3003197	3.94	1.73	2.04	2.63	7.18	5.13	2.76	3.02	4.74				
	NT2RP3003203	10.74	6.48	7.57	9.78	9.35	10.34	12.74	12.49	16.29			*	+
	NT2RP3003204	5.10	4.07	4.28	9.44	9.51	9.35	6.59	6.58	5.8	**	+	*	+
	NT2RP3003210	2.87	2.26	2.76	4.58	4.94	5.68	4.02	4.31	4.86	**	+	**	+
	NT2RP3003212	3.99	3.41	3.08	11.16	9.44	5.92	5.65	5.21	4.76	*	+	**	+
15	NT2RP3003213	3.64	1.51	1.06	6.12	6.44	4.09	4.51	3.54	3.74	*	+		
	NT2RP3003224	4.97	2.24	2.03	5.15	4.35	3.48	1.88	2.89	5.66				
	NT2RP3003226	6.57	4.20	3.82	5.03	7.40	7.29	3.35	4.03	3.53				
	NT2RP3003230	5.88	2.80	3.00	5.34	6.53	3.95	6.24	6.52	4.98				
	NT2RP3003235	5.68	3.50	3.55	11.57	10.99	8.51	10.86	10.22	8.85	**	+	**	+
20	NT2RP3003242	2.60	1.56	1.56	2.17	2.65	0.82	2.88	3.62	2.34				
	NT2RP3003251	6.96	4.06	5.58	8.26	9.86	10.16	5.03	5.10	5.01	*	+		
	NT2RP3003252	3.92	3.17	2.70	4.36	6.32	3.73	3.3	3.53	3.19				
	NT2RP3003258	4.44	4.88	5.51	5.73	7.67	6.20	6.76	5.52	8.07				
	NT2RP3003260	10.73	5.21	4.49	5.79	7.69	5.80	4.33	3.45	7.99				
	NT2RP3003264	3.02	3.32	2.19	15.38	18.88	12.82	6.5	5.90	7.82	**	+	**	+
25	NT2RP3003273	3.18	1.91	3.15	2.64	2.58	3.24	1.86	3.56	1.93				
	NT2RP3003278	3.16	1.06	0.85	1.38	1.88	2.32	0.32	2.37	2.1				
	NT2RP3003280	11.26	9.07	8.30	12.96	14.31	12.01	8.63	10.92	9.88	*	+		
	NT2RP3003282	2.12	1.63	1.57	3.75	3.52	2.64	2.53	3.71	3.58	*	+	*	+
	NT2RP3003290	6.74	3.39	5.29	8.39	9.77	12.47	5.55	7.58	4.52	*	+		
30	NT2RP3003301	3.39	1.66	2.31	5.80	5.15	3.88	3.51	3.63	2.51	*	+		
	NT2RP3003302	4.39	1.94	0.70	3.91	4.34	3.52	1.87	2.40	2.1				
	NT2RP3003311	6.06	3.51	2.81	1.70	1.60	1.58	1.38	2.35	2.23				
	NT2RP3003312	2.65	1.71	1.08	1.61	2.31	2.14	2.34	3.94	2.4				
	NT2RP3003313	2.10	1.55	1.28	2.78	3.32	3.29	2.46	3.52	2.12	**	+		
35	NT2RP3003327	4.75	3.06	2.77	5.48	4.57	3.91	2.76	4.36	2.87				
	NT2RP3003330	2.85	1.28	1.93	2.62	3.38	1.73	2.22	3.78	2.76				
	NT2RP3003344	2.79	2.00	1.76	2.66	2.98	3.04	2.26	1.95	1.8				
	NT2RP3003346	5.06	3.51	3.24	6.69	7.03	5.74	4.23	5.12	4.21	*	+		
	NT2RP3003349	9.03	3.41	4.20	7.42	11.99	8.27	4.03	4.39	5.81				
40	NT2RP3003353	2.34	1.65	0.86	3.37	3.35	2.15	1.51	2.11	2.73				
	NT2RP3003354	28.51	16.58	19.06	32.92	34.54	31.72	24.06	25.56	26.43	*	+		
	NT2RP3003368	4.73	3.35	3.40	3.00	5.12	6.89	5.78	4.93	4.85				
	NT2RP3003375	7.10	4.96	7.12	8.55	8.55	5.98	2.32	4.29	4.97				
	NT2RP3003377	7.20	4.93	4.97	2.66	4.68	3.75	3.7	3.85	3.56				
	NT2RP3003384	2.46	2.07	1.01	3.30	3.65	2.66	3.02	2.86	2.88				
45	NT2RP3003385	5.42	4.79	5.32	4.48	4.42	6.30	6.9	4.85	5.01				
	NT2RP3003396	9.36	4.73	3.86	5.45	9.23	6.23	6.71	5.33	6.72				
	NT2RP3003403	3.05	1.65	1.51	5.41	4.67	5.69	2.27	2.49	2.78	**	+		
	NT2RP3003409	2.84	1.35	2.12	3.28	3.13	1.88	2.79	2.04	3.14				
	NT2RP3003411	8.55	4.92	6.03	7.49	10.91	12.20	6.24	6.43	9.99				
50	NT2RP3003420	4.15	2.44	2.36	6.31	7.10	8.42	3.61	4.27	6.2	**	+		
	NT2RP3003425	3.63	2.52	1.95	2.10	3.46	2.62	1.83	3.37	4.24				
	NT2RP3003426	9.31	6.11	5.45	12.48	10.53	11.34	9.4	9.12	8.46	*	+		
	NT2RP3003427	8.99	4.74	5.99	5.37	7.25	6.52	7.72	5.96	8.43				
	NT2RP3003433	9.63	4.28	3.87	11.80	10.94	8.04	5.78	4.80	7.07				
55	NT2RP3003437	18.34	7.27	6.59	15.83	19.21	15.84	12.02	13.41	15.15				
	NT2RP3003448	6.95	4.49	2.68	9.13	8.95	5.55	5.56	5.02	6.11				
	NT2RP3003455	8.08	3.54	2.51	10.14	9.73	10.02	5.84	8.37	5.46	*	+		

Table 277

	NT2RP3002823	1.32	1.08	1.04	1.83	2.17	1.81	1.57	3.57	2.5	**	+		
	NT2RP3002825	7.13	4.05	4.87	6.63	6.04	8.47	4.09	5.57	4.15				
5	NT2RP3002829	3.03	2.45	2.63	5.74	5.50	4.90	3	3.82	3.79	**	+		
	NT2RP3002831	3.87	3.21	2.77	3.69	2.99	3.89	2.66	2.74	2.29				
	NT2RP3002836	14.03	6.74	6.74	9.92	15.02	8.10	13.6	10.55	13.13				
	NT2RP3002845	6.06	2.27	2.32	3.35	4.67	5.99	2.22	2.92	5.24				
	NT2RP3002852	2.14	1.57	1.15	1.52	1.72	1.72	1.78	2.42	2.44				
10	NT2RP3002861	4.05	2.12	1.50	1.55	2.01	4.44	1.39	3.44	3.12				
	NT2RP3002869	6.92	5.64	4.79	4.48	4.94	3.03	3.48	5.21	5.99				
	NT2RP3002874	3.62	2.41	3.09	2.41	2.83	2.25	3.7	5.14	4.58				
	NT2RP3002876	6.38	5.46	5.19	8.34	12.34	10.89	6.16	7.19	7.18	*	+		
	NT2RP3002877	4.36	2.55	2.24	6.28	5.72	7.39	4.17	3.78	4.69	*	+		
15	NT2RP3002887	2.31	2.06	1.28	2.41	6.33	3.71	2.23	1.91	2.99				
	NT2RP3002900	4.62	3.12	1.94	6.79	7.22	4.89	6.77	4.56	5.42	*	+		
	NT2RP3002902	13.48	7.11	7.49	17.13	16.57	10.16	8.66	6.18	6.66				
	NT2RP3002909	33.33	17.88	18.92	24.91	27.67	27.33	23.19	23.81	25.55				
	NT2RP3002911	2.05	1.51	2.25	2.06	2.34	3.42	1.9	2.88	2.46				
20	NT2RP3002948	2.87	2.05	2.73	3.15	3.80	3.22	3.02	3.24	4.14				
	NT2RP3002953	2.95	2.20	2.80	3.91	2.99	2.13	3.94	4.99	3.35				
	NT2RP3002955	3.21	2.28	2.19	2.68	3.66	2.17	2.8	4.04	3.2				
	NT2RP3002958	5.15	1.89	1.75	8.65	9.49	5.11	5.86	5.70	7.9				
	NT2RP3002969	8.37	4.79	4.07	7.09	7.89	5.99	3.82	5.59	8.02				
25	NT2RP3002972	2.45	1.77	1.17	3.30	4.53	6.41	2.37	3.50	4.2	*	+		
	NT2RP3002978	3.51	1.12	0.76	1.57	2.29	1.16	1.76	2.22	2.49				
	NT2RP3002983	2.09	1.72	1.47	2.93	4.10	4.53	1.5	4.04	1.42	*	+		
	NT2RP3002985	2.93	1.24	0.64	1.80	1.57	1.56	1.03	3.24	1.64				
	NT2RP3002988	3.04	1.50	1.33	2.69	2.87	2.12	2.09	2.69	1.72				
30	NT2RP3003000	5.52	4.04	3.47	8.75	7.05	6.47	5.37	5.35	7.11	*	+		
	NT2RP3003008	3.30	1.49	1.41	3.13	2.40	2.15	3.61	1.58	2.05				
	NT2RP3003012	5.75	2.52	2.34	2.71	2.38	1.98	3.89	1.73	1.65				
	NT2RP3003015	3.67	2.39	1.41	2.11	1.98	2.12	2.64	2.73	1.76				
	NT2RP3003018	5.19	3.49	2.94	3.09	5.88	7.34	2.45	3.41	8.68				
35	NT2RP3003028	4.42	2.89	2.76	3.64	5.83	5.34	3.92	2.05	3.21				
	NT2RP3003029	5.92	3.71	3.59	6.44	6.11	4.11	7.41	7.78	5.42				
	NT2RP3003032	8.58	6.19	7.17	18.73	18.81	11.60	10.2	11.99	14.12	*	+	*	+
	NT2RP3003041	0.23	0.21	0.07	0.41	0.42	0.07	0.35	0.34	-0.17				
	NT2RP3003044	7.25	3.53	3.53	7.47	6.31	4.80	5.47	4.15	4.63				
40	NT2RP3003047	14.58	8.48	8.68	11.39	12.06	11.40	11.77	9.28	11.88				
	NT2RP3003050	6.53	2.71	3.77	5.22	5.47	3.84	5.66	4.93	4.39				
	NT2RP3003053	17.07	9.71	8.94	14.88	15.92	20.90	14.19	12.88	11.32				
45	NT2RP3003059	2.32	1.74	2.11	2.95	2.30	1.48	1.32	1.45	1.42		*	-	
	NT2RP3003061	4.13	2.99	2.62	3.51	4.22	2.44	3.64	4.14	3.12				
	NT2RP3003068	7.07	5.01	4.05	8.08	8.01	6.86	3.94	4.27	5.35				
	NT2RP3003071	7.18	5.69	5.64	19.53	14.10	9.02	3.53	6.99	4.86				
50	NT2RP3003076	20.24	13.69	11.73	17.25	17.10	20.23	12.75	12.44	19.06				
	NT2RP3003078	6.31	1.99	2.60	4.81	6.42	5.61	4.7	3.16	4.19				
	NT2RP3003081	5.58	3.59	4.40	7.90	10.09	9.19	6.17	5.95	6.74	**	+	*	+
	NT2RP3003090	4.22	2.78	2.81	6.19	7.29	6.41	3.45	3.07	3.62	**	+		
	NT2RP3003097	2.80	1.80	2.13	3.12	4.85	3.19	4.18	3.63	2.96		*	+	
55	NT2RP3003098	3.43	1.98	2.02	2.28	3.12	2.15	2.19	2.67	2.43				
	NT2RP3003101	5.48	5.07	5.35	6.08	7.76	5.95	4.99	7.09	5.03				
	NT2RP3003109	14.31	7.48	6.90	18.37	16.28	14.44	15.14	17.70	14.3				
	NT2RP3003121	150.76	6.07	4.19	6.53	16.16	5.37	32.59	5.96	252				
	NT2RP3003133	6.04	4.14	3.20	8.62	13.38	13.91	5.16	4.56	8.91	*	+		
	NT2RP3003137	10.77	5.97	6.19	4.43	6.11	4.14	3.41	3.49	4.4				
	NT2RP3003138	5.81	4.35	3.40	6.66	5.96	5.22	1.99	2.76	2.93				

Table 276

	NT2RP3002582	12.31	7.23	7.62	9.16	12.52	12.19	7.07	6.55	8.27				
5	NT2RP3002587	2.59	1.37	0.54	2.46	2.67	3.02	1.24	1.89	1.22				
	NT2RP3002590	10.29	5.66	7.55	5.34	4.92	3.70	2.44	4.30	2.27		*	-	
	NT2RP3002602	2.82	1.08	1.45	3.79	2.37	2.51	2.16	2.20	1.92				
	NT2RP3002603	23.80	12.85	10.83	16.77	16.77	18.88	33.04	20.98	28.78				
	NT2RP3002621	5.83	2.17	2.11	2.73	3.73	3.84	3.77	3.43	4.67				
10	NT2RP3002622	6.46	4.71	3.37	7.18	6.32	5.80	5.41	4.46	6.55				
	NT2RP3002624	1.38	1.46	0.86	2.16	2.27	1.71	1.92	2.31	2.23	*	+	*	+
	NT2RP3002628	3.88	4.12	4.54	3.93	5.95	4.39	6.01	5.25	6.35		*	*	+
	NT2RP3002629	17.56	11.86	13.81	23.77	21.74	24.60	15.1	15.62	16.2	**	+		
	NT2RP3002631	0.65	0.54	0.71	0.74	2.00	0.23	0.47	2.10	1.77				
15	NT2RP3002647	6.35	4.67	4.32	5.81	4.61	3.54	2.45	3.29	2.94		*	-	
	NT2RP3002649	13.39	5.95	5.65	10.41	9.34	8.49	5.95	5.93	9.13				
	NT2RP3002650	6.81	4.69	4.82	5.81	7.89	6.12	6.83	5.78	9.56				
	NT2RP3002652	5.20	4.74	1.12	4.44	5.82	4.44	3.42	3.65	3.38				
	NT2RP3002654	16.99	10.82	13.04	8.59	8.02	5.74	6.46	6.13	9.06	*	-	*	-
20	NT2RP3002657	6.11	3.63	4.64	10.15	11.45	6.16	9.57	10.27	10.97		**	+	
	NT2RP3002659	1.43	1.66	1.88	2.50	3.07	1.94	1.45	2.43	1.88				
	NT2RP3002660	6.69	4.61	2.72	7.71	9.95	6.32	4.86	5.91	5.04				
	NT2RP3002663	2.95	2.45	2.08	3.55	3.38	2.69	2.33	2.32	1.43				
	NT2RP3002664	4.14	2.04	1.66	3.83	4.46	3.08	3.81	2.61	3.84				
	NT2RP3002667	10.84	11.80	12.31	7.37	13.24	10.35	2.54	3.53	3.86		**	-	
25	NT2RP3002671	4.10	3.38	2.05	3.68	4.13	3.09	3.64	4.14	3.95				
	NT2RP3002682	6.85	6.11	3.50	9.41	10.82	9.25	7.6	6.54	14.33	*	+		
	NT2RP3002684	2.31	2.12	2.06	2.65	2.46	1.95	3.43	3.91	2.52				
	NT2RP3002687	0.81	0.83	0.64	1.63	2.27	2.37	2.18	2.59	1.3	**	+	*	+
	NT2RP3002688	1.90	1.35	1.30	2.68	10.84	4.31	2.62	3.98	4.96		*	+	
30	NT2RP3002698	1.70	1.54	2.28	2.37	1.97	1.69	2.37	4.37	2.27				
	NT2RP3002701	9.13	4.28	3.80	7.31	8.31	6.47	5.76	5.84	9.76				
	NT2RP3002705	21.78	18.18	17.66	50.09	57.33	55.80	17.31	19.57	25.8	**	+		
	NT2RP3002708	8.43	3.13	4.23	10.00	12.33	16.86	6.66	9.06	8.15	*	+		
	NT2RP3002711	10.69	7.85	6.27	14.28	17.41	10.11	7.22	6.34	9.71				
35	NT2RP3002712	75.48	54.09	63.05	72.21	59.93	49.90	55.73	52.68	50.32				
	NT2RP3002713	1.12	1.39	0.99	1.79	1.94	1.51	1.51	1.64	2.24	*	+		
	NT2RP3002721	4.73	3.29	3.45	5.55	8.69	5.41	5.47	5.66	7.4		*	+	
	NT2RP3002722	18.60	15.91	19.67	21.10	20.78	20.71	21.26	14.74	13.19				
	NT2RP3002723	20.89	13.71	12.73	18.65	26.94	25.35	23.58	19.98	24.35				
40	NT2RP3002737	10.83	5.85	5.46	7.36	8.93	8.81	7.12	8.21	8.27				
	NT2RP3002738	3.06	2.31	2.46	3.88	2.93	4.58	4.14	4.86	3.57		*	+	
	NT2RP3002742	78.11	50.55	39.19	56.71	49.99	44.98	24.65	24.79	19.15		*	-	
	NT2RP3002744	1.91	1.57	1.49	3.37	4.81	3.15	4.58	3.73	2.77	*	+	*	+
	NT2RP3002756	2.31	1.24	1.63	1.83	2.14	1.21	1.7	1.60	2.11				
	NT2RP3002757	4.69	3.13	4.35	7.14	8.49	8.18	8.15	8.37	8.37	**	+	**	+
45	NT2RP3002758	7.65	5.42	7.31	13.02	12.93	12.57	12.33	13.43	11.46	**	+	**	+
	NT2RP3002762	17.62	11.52	8.02	10.66	16.28	10.88	8.09	6.08	11.34				
	NT2RP3002763	5.98	3.76	3.67	4.32	6.42	5.16	4.76	6.11	4.92				
	NT2RP3002770	6.69	2.71	1.54	4.12	4.84	3.63	3.88	6.30	6.14				
	NT2RP3002771	4.19	4.34	2.59	8.14	7.86	8.58	10.72	12.24	8.84	**	+	**	+
50	NT2RP3002785	3.87	2.70	2.07	1.69	2.61	1.77	0.79	2.12	2.01				
	NT2RP3002790	2.54	1.59	2.82	4.68	4.85	6.90	3.49	4.63	2.59	*	+		
	NT2RP3002799	2.06	0.55	1.55	2.25	2.19	2.80	1.65	2.16	2.21				
	NT2RP3002801	3.39	2.62	3.03	5.62	4.43	4.91	3.26	3.08	2.61	**	+		
	NT2RP3002802	9.76	4.91	4.56	5.83	7.90	5.66	5.83	5.98	7.36				
55	NT2RP3002810	2.05	2.04	1.36	1.95	2.29	2.16	2.36	3.68	3.36		*	+	
	NT2RP3002818	1.54	1.82	1.16	0.90	1.59	1.73	1.13	2.06	1.73				
	NT2RP3002821	17.00	12.39	12.28	12.51	17.54	13.78	7.96	8.86	8.91		*	-	

Table 275

	NT2RP3002309	6.87	4.15	3.66	6.13	6.93	8.34	2.55	3.41	3.91				
5	NT2RP3002311	4.05	2.38	2.34	4.56	2.55	3.21	2.05	2.83	2.86				
	NT2RP3002315	15.94	11.19	15.32	12.31	8.50	11.56	8.23	8.69	10.92		*	-	
	NT2RP3002319	1.73	1.09	1.94	2.53	2.43	3.11	2.93	2.04	2.66	*	+		
	NT2RP3002324	9.27	3.66	3.72	5.93	9.44	5.66	4.2	5.07	4.43				
	NT2RP3002330	9.95	5.32	3.76	4.42	7.75	7.05	6.63	6.18	5.42				
10	NT2RP3002333	17.93	13.63	12.33	10.81	13.83	11.53	26.44	20.51	21.61		*	+	
	NT2RP3002337	2.63	1.45	1.52	1.90	1.94	2.01	1.38	3.21	2.65				
	NT2RP3002342	15.59	10.64	11.07	10.92	13.50	7.96	9.5	11.72	10.96				
	NT2RP3002343	4.86	3.15	3.42	8.66	7.27	7.64	5.82	6.21	6.54	**	+	*	+
	NT2RP3002351	2.14	1.87	1.48	1.52	1.49	1.39	1.37	2.50	1.29				
15	NT2RP3002352	3.51	2.49	2.09	6.56	3.41	4.41	3.67	4.42	2.26				
	NT2RP3002353	8.54	2.87	2.50	5.68	7.93	6.04	5.65	4.24	3.09				
	NT2RP3002362	10.04	4.71	5.05	6.95	8.81	7.91	8.38	7.04	7.67				
	NT2RP3002363	5.45	3.22	2.99	4.20	6.31	4.65	3.29	3.42	4.78				
	NT2RP3002377	6.53	3.54	3.81	6.50	6.48	4.79	3.11	4.43	2.57				
20	NT2RP3002377	16.05	6.92	7.02	15.78	13.73	11.15	9.35	6.37	9.19				
	NT2RP3002394	3.83	2.35	2.55	5.43	6.35	4.75	5.11	5.17	5.17	*	+	**	+
	NT2RP3002397	1.88	2.06	1.00	2.28	2.42	2.35	2.43	3.20	2.26				
	NT2RP3002399	38.89	13.57	16.73	24.89	24.11	20.07	10.95	10.34	11.58				
	NT2RP3002402	14.13	6.06	6.64	3.90	7.46	3.60	5.13	2.47	3.86				
25	NT2RP3002404	2.69	1.41	1.51	4.63	5.57	6.95	5.03	5.62	5.49	**	+	**	+
	NT2RP3002410	16.74	9.36	8.24	14.55	17.40	14.68	7.71	8.16	9.6				
	NT2RP3002411	5.72	3.09	2.66	5.44	3.76	4.39	3.87	3.60	4.64				
	NT2RP3002414	15.70	13.46	15.51	17.50	19.84	20.94	20.31	15.95	17.64	*	+		
	NT2RP3002430	5.62	3.03	3.26	4.15	6.68	5.69	3.6	5.22	5.76				
	NT2RP3002448	3.21	1.91	1.95	4.68	4.12	2.16	3.43	3.57	3.52				
30	NT2RP3002454	5.75	3.63	2.88	8.65	10.72	8.12	4.17	6.41	5.11	*	+		
	NT2RP3002455	5.96	2.60	2.61	5.44	7.86	5.02	4.61	3.98	4.33				
	NT2RP3002456	19.55	5.82	6.70	24.00	22.06	18.49	6.98	7.59	13.81				
	NT2RP3002462	10.35	5.72	4.60	11.65	13.73	9.93	5.45	7.13	8.04				
	NT2RP3002469	4.02	2.04	2.37	7.68	7.85	6.75	5.57	6.12	6.98	**	+	**	+
35	NT2RP3002470	34.16	21.24	23.62	26.50	31.46	31.78	25.6	23.51	18.11				
	NT2RP3002484	4.96	4.07	3.20	7.26	8.04	8.64	6.14	7.06	7.03	**	+	*	+
	NT2RP3002491	2.02	0.31	0.77	1.88	1.82	1.66	1.79	2.17	2.19				
	NT2RP3002494	5.69	5.46	5.09	5.37	5.09	4.28	11.1	14.53	16.58		**	+	
	NT2RP3002497	7.34	2.87	2.34	7.23	5.25	4.45	4.45	4.17	5.52				
40	NT2RP3002500	6.11	2.15	1.67	4.34	5.06	2.16	2.18	2.29	5.42				
	NT2RP3002501	11.25	5.11	3.44	6.23	6.00	5.47	2.88	5.58	5.46				
	NT2RP3002512	7.00	3.26	2.28	5.82	6.08	6.36	2.87	4.61	8.18				
	NT2RP3002529	3.20	3.16	1.84	7.16	9.33	8.45	4.14	4.40	5.49	**	+	*	+
	NT2RP3002533	7.52	4.47	4.21	12.54	12.31	10.84	8.33	13.60	12.28	**	+	*	+
	NT2RP3002539	6.08	4.61	2.98	8.67	11.27	7.39	2.77	5.22	3.99	*	+		
45	NT2RP3002540	2.20	1.79	1.19	3.09	3.15	2.53	2.67	2.99	2.88	*	+	*	+
	NT2RP3002543	14.24	6.52	5.35	11.36	10.19	12.56	10.27	10.96	8.43				
	NT2RP3002545	4.03	2.04	1.37	6.55	5.22	5.90	5.61	3.59	2.71	*	+		
	NT2RP3002549	2.56	1.25	0.83	5.75	4.78	6.90	5.63	3.95	4.81	**	+	*	+
	NT2RP3002552	2.93	2.06	2.41	3.32	5.85	3.49	4.06	3.68	4.2		**	+	
50	NT2RP3002558	7.05	4.19	4.48	9.57	11.91	11.02	10.69	7.40	9.14	**	+	*	+
	NT2RP3002565	4.40	2.70	2.23	5.52	4.89	4.10	2.94	2.79	3.23				
	NT2RP3002566	4.15	3.12	3.18	4.65	4.50	3.46	4.21	2.25	2.13				
	NT2RP3002571	1.43	0.64	1.11	2.38	2.79	1.21	2	1.13	1.01				
	NT2RP3002572	5.68	2.77	2.24	4.20	4.73	4.39	4.25	2.73	1.87				
55	NT2RP3002573	12.53	5.63	5.03	10.21	9.69	14.18	6.47	7.06	5.93				
	NT2RP3002577	16.44	10.30	7.27	9.02	16.56	19.10	11.76	11.75	13.42				
	NT2RP3002579	5.14	1.77	2.75	2.43	7.06	4.13	4.98	5.48	3.32				

Table 274

	NT2RP3002064	5.17	3.05	2.46	4.06	7.44	4.88	4.84	4.54	4.14				
5	NT2RP3002071	2.33	1.51	1.99	1.86	2.27	1.50	2.16	2.61	2.43				
	NT2RP3002073	5.31	4.25	4.41	3.45	4.48	3.77	3.46	5.18	3.88				
	NT2RP3002074	3.99	3.21	3.54	3.26	5.35	3.47	3.41	4.15	2.51				
	NT2RP3002075	4.75	2.10	2.19	6.52	7.60	4.15	6.03	5.22	5				
	NT2RP3002077	8.02	3.34	2.61	6.63	4.07	3.18	5.14	4.74	2.68				
10	NT2RP3002081	10.07	7.99	7.00	4.79	4.27	3.26	2.76	2.42	1.41	*	-	**	-
	NT2RP3002086	4.94	3.90	3.43	7.01	9.40	7.91	6.79	5.61	5.45	**	+	*	+
	NT2RP3002094	55.21	38.13	49.40	26.53	35.64	30.76	29.38	24.30	29.05	*	-	*	-
	NT2RP3002096	2.03	2.45	2.09	2.34	2.63	1.70	2.31	1.94	2.22				
	NT2RP3002097	4.81	2.56	2.66	7.07	9.45	4.39	4.28	5.92	5.09				
15	NT2RP3002098	1.30	1.49	2.04	3.02	3.52	2.23	1.86	1.80	1.76	*	+		
	NT2RP3002102	4.48	2.97	2.73	5.04	5.32	5.08	5.06	4.28	4.93	*	+		
	NT2RP3002106	5.41	2.39	2.38	9.26	7.89	8.90	6.1	3.83	3.57	**	+		
	NT2RP3002108	6.53	3.49	4.50	3.88	5.75	3.58	3.09	4.07	3.18				
	NT2RP3002109	11.23	5.02	4.28	16.19	18.27	13.88	14.35	12.01	12.31	*	+		
20	NT2RP3002110	23.37	14.84	16.48	34.91	29.71	40.33	23.01	21.75	23.48	*	+		
	NT2RP3002113	11.63	9.01	7.67	6.51	7.35	7.47	7.32	7.10	6.45				
	NT2RP3002120	1.55	1.48	1.08	2.91	3.24	1.92	2.33	3.13	2.18	*	+	*	+
	NT2RP3002121	3.47	2.28	2.84	4.15	6.05	2.79	2.22	3.39	2.01				
	NT2RP3002126	11.23	6.99	4.03	8.17	8.24	7.23	16.66	12.35	16.36		*	+	
25	NT2RP3002128	13.16	6.63	6.22	10.39	10.13	7.09	9.73	7.03	10.29				
	NT2RP3002130	7.94	5.84	4.52	8.35	9.12	8.25	8.69	6.14	9.87				
	NT2RP3002133	7.00	4.13	2.94	10.10	13.02	11.57	10.36	9.95	10.86	**	+	**	+
	NT2RP3002136	10.87	7.59	6.07	13.09	20.57	19.22	14.35	15.02	15.43	*	+	**	+
	NT2RP3002140	4.41	4.46	5.24	5.99	5.61	7.54	7.49	4.80	5.22				
	NT2RP3002142	7.81	6.29	3.94	14.63	15.34	11.73	11.3	15.25	13.24	**	+	*	+
30	NT2RP3002146	7.61	4.78	4.77	10.91	13.18	6.97	4.8	6.21	4.65				
	NT2RP3002147	22.06	11.75	12.01	9.65	10.83	10.56	11.86	8.17	9.38				
	NT2RP3002151	14.60	11.05	8.77	13.96	13.74	12.27	8.15	8.64	12.04				
	NT2RP3002155	8.16	6.32	4.96	8.79	7.65	4.96	6.19	7.55	7.22				
	NT2RP3002156	2.21	1.36	0.96	3.23	3.14	2.36	3.21	3.07	3.25	*	+	*	+
35	NT2RP3002160	3.98	3.19	1.94	3.32	4.52	5.20	4.3	1.89	4.12				
	NT2RP3002163	18.81	11.61	12.16	18.87	21.42	15.74	12.51	9.05	10.05				
	NT2RP3002165	6.12	5.16	5.75	6.38	8.10	3.82	6.23	5.63	7.23				
	NT2RP3002166	5.72	3.53	1.35	2.95	5.16	3.30	2.3	3.24	3.17				
	NT2RP3002173	5.34	3.03	2.78	9.80	6.20	7.21	5.06	5.00	4.94	*	+		
40	NT2RP3002174	5.68	2.49	1.67	7.29	8.21	9.12	9.02	7.21	12.43	*	+	*	+
	NT2RP3002181	9.68	7.50	5.24	4.48	4.92	3.59	2.61	2.36	2.48		*	-	
	NT2RP3002185	3.81	2.37	1.77	2.88	7.87	3.22	3.57	3.44	2.54				
	NT2RP3002193	7.51	6.09	4.76	5.28	9.69	7.23	6.2	5.26	7.9				
	NT2RP3002204	2.89	2.47	0.95	9.64	8.53	14.75	4.05	4.67	4.6	*	+	*	+
	NT2RP3002244	4.56	5.32	5.18	4.63	6.32	6.34	4.51	3.44	3.59		*	-	
45	NT2RP3002248	8.18	5.72	5.54	14.10	16.32	12.91	11.02	10.26	11.54	**	+	**	+
	NT2RP3002253	6.83	4.26	3.08	6.54	5.65	6.66	3.58	3.16	4.05				
	NT2RP3002255	44.02	22.63	19.64	26.45	26.34	31.64	17.22	13.77	17.68				
	NT2RP3002264	5.83	3.17	2.53	6.13	7.07	6.24	4.47	6.97	4.95				
	NT2RP3002267	4.61	2.60	2.31	3.48	4.99	3.73	3.57	2.66	3.09				
50	NT2RP3002273	14.02	8.03	6.96	15.74	16.07	14.50	9.58	10.63	9.37				
	NT2RP3002276	5.72	2.96	3.52	5.50	5.94	5.34	3.99	5.68	5.16				
	NT2RP3002281	7.91	5.75	6.50	6.21	6.83	6.47	4.32	5.43	5.21				
	NT2RP3002286	2.46	1.62	2.05	3.65	3.52	2.26	2.34	3.39	3.14				
	NT2RP3002297	56.91	27.98	24.70	67.63	63.96	45.16	26.65	22.90	25.3				
55	NT2RP3002301	9.96	5.96	5.15	5.72	8.90	9.72	8.36	7.88	9.26				
	NT2RP3002303	10.45	6.01	4.55	8.24	9.49	7.38	8.89	7.75	8.68				
	NT2RP3002304	1.01	1.07	1.38	3.55	2.86	2.06	2.84	4.66	2.09	*	+		

Table 273

	NT2RP3001771	3.51	2.93	3.35	3.89	4.06	3.55	3.61	4.21	5.23			
5	NT2RP3001777	4.09	2.96	3.01	5.51	4.45	3.91	4.86	5.16	6		*	+
	NT2RP3001782	2.53	2.57	1.95	6.76	6.36	6.69	4.29	4.57	3.41	**	+	*
	NT2RP3001792	5.75	4.70	5.90	6.11	8.15	9.14	6.11	4.96	5.99			
	NT2RP3001799	4.41	4.21	3.75	7.39	9.01	7.29	5.88	7.01	5.73	**	+	**
	NT2RP3001819	6.61	3.33	1.74	4.45	5.18	4.58	4.38	3.34	4.47			
10	NT2RP3001829	60.87	38.63	36.73	56.07	52.70	55.16	28.32	28.08	35.16			
	NT2RP3001836	10.17	5.74	4.77	10.85	13.55	11.18	6.57	5.69	7.14			
	NT2RP3001839	15.46	12.06	10.35	17.55	22.87	17.91	17.89	15.53	21.32	*	+	
	NT2RP3001844	5.39	4.22	4.08	8.68	8.00	8.70	4.83	4.18	5.54	**	+	
	NT2RP3001848	8.51	3.03	3.37	7.54	6.39	7.94	7.05	8.18	5.83			
15	NT2RP3001854	4.31	3.66	2.93	4.93	7.64	5.42	5.84	9.19	10.46		*	+
	NT2RP3001855	1.08	0.62	0.41	0.88	3.15	1.50	2.17	1.51	1.24			
	NT2RP3001857	8.74	5.14	3.23	3.88	5.79	4.95	4.34	4.47	3.21			
	NT2RP3001858	5.96	2.68	3.12	1.87	2.69	2.83	2.52	3.04	2.59			
	NT2RP3001861	8.95	6.91	5.65	7.71	8.95	8.02	9.41	9.63	9.39			
	NT2RP3001866	1.78	1.67	1.30	2.40	3.59	1.96	3.62	3.94	3.33		**	+
20	NT2RP3001871	1.22	1.47	1.24	4.28	5.33	4.06	5.94	5.76	6.13	**	+	**
	NT2RP3001874	2.39	1.48	1.04	1.60	1.73	1.49	2.15	3.07	2.44			
	NT2RP3001878	1.89	1.50	2.48	4.52	7.04	3.00	1.74	2.47	2.05			
	NT2RP3001885	4.23	3.76	3.61	4.08	6.00	8.45	4.94	5.08	4.08			
	NT2RP3001896	3.95	2.31	1.26	4.38	7.80	4.28	4.49	2.83	4.64			
25	NT2RP3001898	12.61	5.06	3.64	6.11	6.18	5.92	8.68	7.13	11.31			
	NT2RP3001899	5.05	3.28	2.34	3.69	5.19	3.08	2.74	3.58	3.91			
	NT2RP3001901	12.98	8.89	8.12	8.50	8.51	10.47	8.45	6.54	7.26			
	NT2RP3001915	6.53	3.55	4.50	3.73	7.04	4.19	2.46	3.27	3.28			
	NT2RP3001926	0.32	0.45	0.32	1.03	1.16	1.31	0.6	2.68	0.45	**	+	
30	NT2RP3001929	2.79	2.04	3.11	3.82	2.97	3.77	2.42	3.15	2.72			
	NT2RP3001931	4.35	3.16	3.68	6.47	4.72	7.93	3.59	3.28	4.34			
	NT2RP3001938	7.26	2.97	4.06	7.92	6.46	6.68	4	4.10	3.17			
	NT2RP3001943	14.11	5.27	4.51	10.79	10.92	8.33	5.43	5.45	5.13			
	NT2RP3001944	3.45	2.33	1.32	2.72	2.97	3.31	3.63	3.49	2.49			
35	NT2RP3001945	7.29	7.10	5.59	8.17	9.64	11.51	6.42	7.34	6.69	*	+	
	NT2RP3001947	4.79	4.51	3.45	5.88	6.32	6.85	5.07	6.05	6.08	*	+	*
	NT2RP3001949	2.69	1.52	2.67	4.00	3.55	3.46	2.68	2.84	2.52	*	+	
	NT2RP3001952	16.48	13.65	16.67	12.37	9.06	10.48	18.01	17.39	16.21	*	-	
	NT2RP3001954	5.28	2.86	2.85	5.44	4.55	3.42	3.76	3.67	4.11			
40	NT2RP3001956	34.22	13.29	14.18	28.43	28.08	22.94	14.79	12.62	14.22			
	NT2RP3001967	7.52	2.65	2.30	9.80	9.24	5.06	8.63	5.51	4.88			
	NT2RP3001969	7.99	4.86	4.65	5.70	7.31	4.72	3.47	2.46	4.31			
	NT2RP3001976	7.58	3.71	3.57	8.43	12.72	10.69	5.69	4.81	4.65	*	+	
	NT2RP3001986	4.77	4.42	3.72	5.84	6.16	3.49	3.93	4.27	4.43			
	NT2RP3001989	0.59	0.37	0.61	1.26	1.01	1.46	1.37	2.34	1.2	**	+	*
45	NT2RP3002002	4.58	2.14	1.97	6.96	7.70	8.62	3.16	5.19	5.28	**	+	
	NT2RP3002004	2.02	1.54	1.44	3.44	3.14	2.45	2.24	2.01	2.64	*	+	
	NT2RP3002007	2.30	1.16	1.11	2.63	4.31	2.50	1.57	1.64	1.85			
	NT2RP3002014	4.46	3.07	2.32	5.12	6.41	4.59	6.11	3.83	4.25			
	NT2RP3002015	7.60	4.06	4.17	6.58	5.55	3.85	6.25	4.00	5.51			
50	NT2RP3002033	1.85	1.50	1.64	2.80	2.86	2.23	1.62	2.56	2.12	*	+	
	NT2RP3002045	1.82	1.00	1.37	1.94	4.75	2.88	1.69	2.27	2.09			
	NT2RP3002054	2.00	1.59	0.94	1.90	1.75	2.02	1.61	2.56	2.14			
	NT2RP3002056	2.28	1.93	1.78	6.00	7.33	7.83	2.85	2.69	4.61	**	+	
	NT2RP3002057	1.99	1.12	1.41	3.14	2.25	1.70	1.48	2.86	1.82			
55	NT2RP3002061	16.71	9.57	7.36	24.61	19.84	16.31	10.52	9.51	8.46			
	NT2RP3002062	2.33	1.47	0.86	3.09	3.51	2.69	1.64	2.02	3.16	*	+	
	NT2RP3002063	8.43	3.19	2.56	5.90	5.68	4.65	5.99	6.66	4.47			

Table 272

	NT2RP3001539	12.56	6.40	7.00	10.99	10.52	8.15	6.27	6.65	5.19				
5	NT2RP3001542	3.56	1.19	1.50	6.99	9.11	5.28	2.14	3.06	2.68 *	+			
	NT2RP3001549	9.80	7.45	10.38	11.31	10.30	10.04	7.8	5.81	7.76				
	NT2RP3001554	3.44	2.57	2.68	4.38	5.21	3.74	3.1	4.12	3.42 *	+			
	NT2RP3001560	1.98	0.84	1.82	2.21	1.46	2.33	2.57	1.64	2.81				
	NT2RP3001561	7.62	4.57	4.64	6.91	8.11	8.03	7.34	7.68	6.78				
10	NT2RP3001564	12.59	4.99	5.10	22.94	20.84	14.16	5.83	7.51	11.43 *	+			
	NT2RP3001568	10.68	5.54	6.19	6.22	5.75	5.19	2.58	3.78	3.57				
	NT2RP3001575	10.33	5.99	5.32	11.60	12.09	8.47	6.09	5.98	6.46				
	NT2RP3001580	3.56	1.35	1.99	5.39	3.01	3.50	2.91	3.43	3.66				
	NT2RP3001587	9.27	5.60	6.48	9.67	8.64	7.91	3.57	5.67	3.81				
15	NT2RP3001589	4.49	2.24	2.17	4.59	7.05	6.18	4.42	5.38	3.17 *	+			
	NT2RP3001592	4.37	2.01	2.87	4.75	5.39	5.86	3.63	4.01	2.99 *	+			
	NT2RP3001607	0.30	0.54	0.84	0.71	1.22	1.55	0.82	2.08	0.53				
	NT2RP3001608	7.31	2.87	2.62	6.20	4.67	5.11	3.69	5.29	6.29				
	NT2RP3001613	11.75	4.76	3.72	8.30	8.98	5.57	5.89	6.91	7.14				
20	NT2RP3001619	4.55	2.53	2.20	3.59	4.12	3.29	2.64	4.30	2.99				
	NT2RP3001621	7.09	6.13	3.47	2.20	2.82	2.93	1.51	2.76	2.37		*	-	
	NT2RP3001629	3.07	1.05	1.36	2.67	2.54	2.74	1.29	3.63	1.56				
	NT2RP3001630	4.04	2.39	2.24	3.71	3.71	2.59	1.51	3.51	0.99				
	NT2RP3001631	24.78	10.11	12.40	17.73	20.88	13.17	4.28	8.91	6.44				
	NT2RP3001634	9.27	2.72	5.54	7.96	8.15	7.28	4.29	5.79	4.53				
25	NT2RP3001642	5.13	3.42	2.92	6.54	7.68	6.47	5.19	3.70	3.73 *	+			
	NT2RP3001646	3.27	1.84	0.92	3.18	2.57	2.35	5	2.95	3.44				
	NT2RP3001650	3.62	2.89	1.93	2.64	3.29	4.41	2.44	1.58	2.48				
	NT2RP3001667	1.93	2.07	1.35	2.81	3.65	4.62	4.85	5.42	7.49 *	+	**	+	
	NT2RP3001671	7.66	4.46	4.89	5.72	6.98	5.49	3.11	2.99	4.06				
30	NT2RP3001672	5.04	4.31	3.86	3.93	4.78	3.32	4.59	4.37	7.43				
	NT2RP3001676	3.97	2.04	5.02	4.84	5.72	3.79	2.56	2.60	3.1				
	NT2RP3001678	5.11	3.61	3.12	4.03	3.95	2.98	4.85	3.51	3.88				
	NT2RP3001679	5.80	3.94	3.38	8.40	8.81	5.85	11	8.10	8.4 *	+	*	+	
	NT2RP3001682	11.08	7.03	6.66	4.48	3.93	2.41	1.86	2.18	2.25 *	-	*	-	
35	NT2RP3001685	5.84	2.49	1.45	5.20	7.06	5.72	3.81	3.24	3.24				
	NT2RP3001688	9.98	5.14	4.96	11.67	15.18	13.11	7.75	5.30	4.79 *	+			
	NT2RP3001690	6.37	3.50	2.59	4.35	7.48	8.72	4.02	4.96	4.94				
	NT2RP3001693	13.26	8.38	9.13	9.74	11.97	8.26	6.72	8.53	7.59				
	NT2RP3001696	6.95	4.47	3.30	15.86	17.48	7.56	13.16	12.78	11.08		**	+	
40	NT2RP3001698	6.30	3.93	3.04	7.50	5.16	4.97	10.41	6.02	8.18				
	NT2RP3001708	3.49	1.19	1.37	2.49	3.70	3.38	4.25	2.37	2.33				
	NT2RP3001712	11.74	6.82	5.41	22.86	35.26	39.54	12.07	11.43	15.14 *	+			
	NT2RP3001716	7.22	3.02	4.03	8.79	10.51	6.60	4.73	4.70	5.85				
	NT2RP3001724	15.75	4.14	3.21	5.86	6.17	7.63	4.16	4.41	4.61				
	NT2RP3001727	8.66	6.49	5.38	14.44	7.82	11.73	11.95	13.12	10.93		*	+	
45	NT2RP3001729	1.93	0.96	0.61	2.40	2.57	2.22	2.16	2.35	2.73 *	+	*	+	
	NT2RP3001730	6.71	4.57	7.74	11.66	10.98	8.11	6.76	8.86	5.97				
	NT2RP3001733	2.88	2.06	0.55	2.95	3.43	1.42	2.02	2.52	2.06				
	NT2RP3001737	6.70	4.04	4.02	6.45	5.41	5.38	5.72	3.92	6.08				
	NT2RP3001738	10.91	6.90	7.77	7.27	7.41	7.04	6.92	5.83	6.78				
50	NT2RP3001739	5.34	4.75	4.43	4.78	6.81	5.30	5.03	4.71	6.57				
	NT2RP3001742	5.50	3.13	4.00	3.39	9.70	3.77	4.55	5.25	8.02				
	NT2RP3001751	13.48	12.01	10.94	15.12	15.40	18.57	7.79	9.88	12.42 *	+			
	NT2RP3001752	4.05	3.78	2.59	14.37	14.59	7.40	13.28	13.75	10.73 *	+	**	+	
	NT2RP3001753	4.22	3.12	2.93	5.12	4.27	8.95	2.67	3.47	2.04				
55	NT2RP3001754	24.40	11.37	10.27	18.41	20.20	17.55	14.78	11.55	16.09				
	NT2RP3001756	3.63	3.86	3.16	12.94	21.73	28.36	7.24	4.83	10.97 *	+			
	NT2RP3001764	6.68	4.75	3.99	4.90	5.39	5.66	4.26	4.39	5.91				

Table 271

	NT2RP3001300	6.60	4.50	3.63	5.55	5.25	4.91	6.62	5.73	6.77				
5	NT2RP3001301	4.23	2.95	2.87	6.64	7.54	6.04	6.28	4.21	5.54	**	+		
	NT2RP3001307	3.27	2.88	2.97	3.26	3.20	4.09	4.01	4.56	2.31				
	NT2RP3001310	14.83	12.54	13.73	16.67	19.61	13.33	4.87	4.68	5.5		**	-	
	NT2RP3001318	2.74	0.91	1.95	3.13	4.02	2.55	2.31	3.93	2.1				
	NT2RP3001322	1.63	0.90	0.90	2.74	1.95	2.82	2.79	4.85	2.34	*	+		
10	NT2RP3001325	24.22	12.72	10.92	7.56	8.85	7.79	5.36	4.73	4.91				
	NT2RP3001338	15.76	12.66	9.88	10.48	14.22	14.25	12.53	9.24	15.76				
	NT2RP3001339	4.32	1.49	2.10	2.91	3.84	2.95	3.46	1.89	3.19				
	NT2RP3001340	19.62	12.27	15.41	18.07	21.26	18.46	16.08	15.90	21.63				
	NT2RP3001341	4.04	2.16	2.75	3.64	4.76	3.69	3.08	3.25	2.32				
	NT2RP3001354	12.69	8.27	10.24	14.38	16.19	12.96	8.57	6.12	4.87				
15	NT2RP3001355	3.39	2.67	2.73	4.52	3.86	4.06	3.69	3.97	3.97	*	+	*	+
	NT2RP3001356	2.63	2.41	2.61	3.21	3.25	2.89	2.82	3.46	1.7	*	+		
	NT2RP3001359	5.31	3.10	1.88	3.19	6.05	4.15	4.41	3.34	3.75				
	NT2RP3001364	6.03	3.09	3.48	5.69	5.56	4.55	3.38	5.70	5.8				
20	NT2RP3001373	5.46	3.57	2.36	4.41	5.80	3.94	5.01	3.68	6.3				
	NT2RP3001374	2.93	1.03	1.18	2.06	2.91	2.46	1.54	1.85	1.14				
	NT2RP3001383	6.37	4.77	6.05	9.28	12.56	10.77	4.11	4.30	3.48	**	+	*	-
	NT2RP3001384	4.58	2.86	3.25	5.41	5.38	4.60	5.49	4.04	4.15				
	NT2RP3001388	3.94	3.65	4.40	11.98	17.15	15.81	10.54	11.04	15.23	**	+	**	+
	NT2RP3001392	3.83	1.90	3.17	5.39	4.17	3.84	3.44	3.44	2.66				
25	NT2RP3001396	2.00	1.30	0.75	2.42	4.93	3.82	4.83	3.81	2.6	*	+	*	+
	NT2RP3001398	11.01	6.05	6.28	7.94	10.96	10.36	8.08	7.65	10.79				
	NT2RP3001399	8.19	4.25	5.07	7.54	8.60	8.41	4.97	7.59	6.74				
	NT2RP3001402	2.09	1.57	1.57	3.12	4.36	4.40	2.46	3.16	5.1	**	+		
	NT2RP3001407	9.10	4.59	5.21	13.05	12.91	13.40	7.95	7.65	8.13	**	+		
30	NT2RP3001416	2.87	2.04	3.00	3.89	8.00	5.00	4.89	5.09	4.41		**	+	
	NT2RP3001420	5.16	2.34	2.93	5.77	5.70	6.45	3.3	5.56	7.47	*	+		
	NT2RP3001425	3.64	1.83	2.78	5.54	5.58	5.80	4.28	4.76	3.32	**	+		
	NT2RP3001426	9.63	6.68	3.99	4.77	6.95	7.51	7.14	4.61	7.91				
	NT2RP3001427	4.50	3.40	2.04	4.15	3.27	4.34	2.81	4.38	3.95				
35	NT2RP3001428	4.16	3.58	4.14	7.37	9.48	9.19	4.5	5.96	3.96	**	+		
	NT2RP3001429	2.71	0.65	1.93	11.45	6.48	6.19	4.59	7.62	3.98	*	+	*	+
	NT2RP3001432	3.34	1.56	1.82	4.80	3.24	3.78	1.92	2.57	3.01				
	NT2RP3001439	6.50	4.98	6.18	6.78	9.50	6.94	5.45	6.68	5.8				
	NT2RP3001441	4.58	1.98	2.38	4.38	3.89	3.43	3.38	5.92	9.79				
40	NT2RP3001446	2.76	1.22	2.57	5.62	7.47	6.18	5.44	6.68	4.2	**	+	*	+
	NT2RP3001447	8.22	4.12	2.95	6.40	8.22	5.10	3.65	5.93	6.09				
	NT2RP3001449	4.73	2.05	2.23	6.25	6.19	5.57	6.13	6.05	7.57	*	+	*	+
	NT2RP3001453	6.27	2.66	2.61	7.65	7.63	7.03	4.7	5.93	5.45	*	+		
	NT2RP3001457	5.03	2.53	2.21	3.77	4.85	3.80	3.24	4.94	2.94				
	NT2RP3001459	2.60	1.82	2.24	2.49	3.26	2.21	2.13	3.94	1.79				
45	NT2RP3001463	3.43	2.23	2.76	3.05	4.78	3.63	2.47	3.86	2.66				
	NT2RP3001466	0.65	0.45	0.93	0.79	1.40	1.78	1.01	1.24	0.81				
	NT2RP3001472	5.02	3.77	3.20	8.65	6.87	6.75	5.25	4.56	5.18	*	+		
	NT2RP3001473	16.30	4.98	4.56	9.54	12.17	8.13	7.39	5.93	7.4				
	NT2RP3001479	11.30	7.78	6.68	11.47	10.59	7.30	7.74	6.55	7.95				
50	NT2RP3001490	1.44	1.38	1.23	3.68	2.94	3.11	4.42	3.30	2.91	**	+	**	+
	NT2RP3001492	3.13	2.23	1.38	5.46	5.82	3.49	2.27	3.77	3.59	*	+		
	NT2RP3001495	4.27	2.41	2.48	4.72	5.59	4.95	3.72	4.06	3.66	*	+		
	NT2RP3001497	3.41	1.98	2.83	6.14	5.70	4.65	3.85	3.87	3.68	*	+		
	NT2RP3001501	3.65	1.22	1.98	4.41	3.90	3.76	3.18	3.14	3.33				
55	NT2RP3001527	8.81	6.07	6.17	11.31	10.29	10.39	6.88	6.90	7.25	*	+		
	NT2RP3001529	9.25	3.58	2.90	11.50	12.88	7.44	5.1	3.82	4.38				
	NT2RP3001538	8.31	2.40	2.73	6.50	6.03	5.12	5.17	5.15	4.98				

Table 270

5	NT2RP3001057	8.67	4.03	5.93	19.26	14.18	12.30	8.42	6.94	7.2	*	+		
	NT2RP3001061	5.88	4.01	4.14	7.75	9.70	8.03	5.42	6.19	4.88	*	+		
	NT2RP3001069	9.78	4.93	-5.43	13.99	17.62	14.76	9.74	9.96	12.86	**	+		
	NT2RP3001074	8.31	4.57	4.04	11.86	10.34	7.95	6.59	7.36	7.45				
	NT2RP3001078	5.34	2.26	4.49	9.51	7.77	7.53	5.94	3.60	5.02	*	+		
	NT2RP3001081	3.83	2.45	4.20	6.12	3.89	6.40	3.56	5.22	3.4				
10	NT2RP3001084	5.54	2.82	2.70	2.36	4.10	1.78	2.85	2.45	3.36				
	NT2RP3001095	1.93	1.69	1.44	3.80	3.49	3.25	2.25	2.83	2.47	**	+	*	+
	NT2RP3001096	4.61	2.92	2.43	5.50	5.58	4.69	7.37	7.57	7.11			**	+
	NT2RP3001097	9.61	7.40	9.00	12.56	12.16	11.92	6.67	6.88	9.12	**	+		
	NT2RP3001107	6.04	4.02	3.50	4.89	5.87	4.23	3.8	4.49	5.02				
15	NT2RP3001109	6.26	3.05	4.30	3.18	4.47	2.65	2.28	2.72	1.85				
	NT2RP3001111	4.22	3.38	2.92	4.13	5.15	4.60	4.36	4.69	3.98				
	NT2RP3001112	28.16	25.89	21.28	24.06	17.68	24.85	10.06	10.75	13.22		**		
	NT2RP3001113	1.79	0.99	0.62	1.34	2.23	1.54	1.11	1.24	1.25				
	NT2RP3001115	3.88	1.85	2.25	7.26	3.45	2.57	3.4	3.91	4.67				
20	NT2RP3001116	3.94	1.69	1.56	4.63	3.42	2.74	3.29	4.13	4.33				
	NT2RP3001119	9.02	6.38	5.74	6.52	9.40	7.53	6.04	4.43	6.5				
	NT2RP3001120	11.82	5.87	8.94	18.20	12.33	18.08	8.42	9.14	10.96	*	+		
	NT2RP3001126	3.38	2.35	3.59	5.64	8.45	7.51	8.01	7.65	6.3	*	+	**	+
	NT2RP3001127	1.21	0.67	1.51	2.88	2.70	1.71	4.11	3.13	5.1	*	+	**	+
25	NT2RP3001133	7.23	4.12	5.49	7.95	8.82	7.67	4.57	6.00	4.72				
	NT2RP3001140	2.84	1.04	1.66	3.30	3.99	3.19	1.56	2.10	3.38	*	+		
	NT2RP3001147	7.62	3.19	3.51	4.05	4.82	4.29	0.77	2.87	1.63				
	NT2RP3001150	5.19	1.79	3.13	6.49	3.73	3.77	3.8	3.66	4.52				
	NT2RP3001152	2.12	0.44	0.89	1.69	1.74	1.98	1.83	2.36	2.08				
	NT2RP3001155	6.90	4.51	4.25	3.69	4.69	3.75	1.73	3.87	3.96				
30	NT2RP3001156	2.47	1.68	1.60	2.59	3.59	3.31	2.51	4.84	4.65	*	+		
	NT2RP3001159	12.19	5.40	5.34	9.00	9.95	7.35	6.84	6.11	6.14				
	NT2RP3001170	7.10	4.60	5.72	9.66	13.09	10.69	5.5	6.89	3.71	*	+		
	NT2RP3001176	9.51	3.49	2.75	17.93	12.62	10.20	6.88	5.97	13.3				
	NT2RP3001195	6.18	2.83	2.96	6.39	10.42	3.54	4.18	5.32	5.17				
35	NT2RP3001209	29.33	14.29	10.79	23.50	28.08	21.04	16.75	19.48	15.61				
	NT2RP3001214	6.63	3.46	3.32	9.82	10.42	9.38	3.48	5.63	3.56	**	+		
	NT2RP3001216	4.48	3.19	3.11	7.11	8.39	8.87	2.58	5.22	3.57	**	+		
	NT2RP3001221	1.19	0.31	0.47	1.55	1.56	1.10	1.01	2.22	0.86				
	NT2RP3001226	7.00	2.58	2.80	4.50	5.21	4.34	3.95	5.75	3.9				
40	NT2RP3001230	2.86	1.59	1.71	4.14	3.19	2.63	1.59	3.61	2.99				
	NT2RP3001232	4.81	1.38	0.57	1.61	2.09	1.97	2.63	1.53	0.99				
	NT2RP3001236	1.71	1.43	0.80	2.59	2.82	2.72	3.58	2.05	2.31	**	+		
	NT2RP3001239	2.21	1.46	1.67	2.79	2.29	1.43	3.36	2.12	2				
	NT2RP3001240	2.39	2.60	2.79	4.11	6.20	4.44	7.84	6.72	4.74	*	+	*	+
45	NT2RP3001245	3.14	1.64	2.84	6.19	9.37	6.48	4.16	3.07	4.85	*	+		
	NT2RP3001253	4.00	1.90	2.62	6.61	7.24	6.92	3.25	4.04	5.99	**	+		
	NT2RP3001259	10.11	5.52	6.66	9.63	10.72	9.87	6.94	7.54	9.1				
	NT2RP3001260	1.75	0.60	0.84	2.44	2.65	2.56	1.25	1.75	2.02	*	+		
	NT2RP3001264	3.80	0.98	1.35	3.72	2.40	2.94	2.21	1.54	2.06				
	NT2RP3001268	5.50	3.38	4.02	7.85	8.76	7.64	4.87	3.66	4.6	**	+		
50	NT2RP3001271	28.62	19.09	17.03	21.24	19.12	21.60	21.92	16.59	24.45				
	NT2RP3001272	5.76	3.32	1.84	5.66	6.83	7.58	3.78	6.70	4.51				
	NT2RP3001274	19.11	14.57	13.97	21.86	23.69	19.32	19.59	16.07	21.69	*	+		
	NT2RP3001275	3.98	2.12	2.06	4.08	3.88	3.61	4.57	5.00	3.17				
	NT2RP3001280	5.95	4.26	3.61	5.15	6.58	6.13	4	4.48	3.31				
55	NT2RP3001281	4.63	3.14	4.04	6.78	5.25	8.51	3.4	3.74	3.77	*	+		
	NT2RP3001288	14.66	10.02	11.01	19.91	17.12	14.80	31.14	30.59	36.12		**	+	
	NT2RP3001297	4.65	2.39	2.87	6.59	5.46	6.16	4.33	3.73	6.69	*	+		

Table 269

	NT2RP3000735	2.00	1.20	0.61	2.06	0.92	1.03	2.17	1.47	1.63			
	NT2RP3000736	3.46	3.21	3.33	4.48	4.58	3.34	3.43	2.28	2.96			
5	NT2RP3000739	15.24	8.34	8.12	11.53	11.36	10.77	13.58	12.81	14.45			
	NT2RP3000742	15.14	9.63	9.98	14.05	14.60	13.15	13.09	11.17	13.06			
	NT2RP3000753	4.09	1.46	2.26	4.87	6.45	3.41	1.81	3.35	5.41			
	NT2RP3000759	4.36	3.02	3.28	9.27	10.72	9.10	9.4	9.92	12.65	**	+	**
	NT2RP3000789	6.97	3.15	3.19	2.62	3.38	3.33	2.9	2.77	2.91			
10	NT2RP3000815	3.08	1.87	2.78	5.08	5.91	5.79	4.34	3.06	3.33	**	+	
	NT2RP3000818	7.88	5.88	4.83	9.79	13.01	13.93	8.4	7.38	10.56	*	+	
	NT2RP3000820	6.70	4.35	2.57	15.50	20.24	18.97	5.35	5.01	5.38	**	+	
	NT2RP3000821	6.58	4.20	3.95	5.67	6.08	4.63	5.13	4.56	4.66			
	NT2RP3000825	0.66	0.26	0.38	1.28	1.09	2.20	0.44	1.29	0.44	*	+	
15	NT2RP3000826	14.31	7.15	8.00	20.59	14.43	14.08	24	29.57	29.39		**	+
	NT2RP3000836	8.67	4.78	5.47	15.61	15.21	9.41	7.61	8.53	8.85	*	+	
	NT2RP3000838	69.68	35.31	38.08	62.74	50.92	57.55	114.4	92.67	110.6		*	+
	NT2RP3000839	3.11	1.70	2.32	2.00	3.56	1.87	3.03	1.30	2.5			
	NT2RP3000841	4.62	3.46	2.85	4.30	8.16	5.93	4.11	3.68	3.13			
20	NT2RP3000845	4.22	3.31	3.16	4.56	7.12	4.56	4.69	3.53	11.01			
	NT2RP3000847	8.01	5.03	4.67	11.17	12.10	10.61	8.29	6.56	5.96	**	+	
	NT2RP3000848	4.58	2.34	3.27	5.39	6.00	5.09	3.72	3.05	5.42	*	+	
	NT2RP3000850	7.12	3.32	4.95	11.87	12.25	13.21	7.48	7.20	7.92	**	+	
	NT2RP3000852	2.41	2.02	3.14	2.50	3.10	2.98	1.15	2.04	2			
25	NT2RP3000859	11.57	6.45	2.66	9.86	9.35	7.35	6.51	5.86	6.19			
	NT2RP3000861	12.29	5.70	6.74	20.57	26.68	20.53	8.96	8.46	14.99	**	+	
	NT2RP3000862	10.74	6.85	6.61	6.87	7.71	5.23	6.09	5.39	7.24			
	NT2RP3000865	2.61	2.77	1.86	4.46	4.70	3.49	3.05	2.82	3.22	*	+	
	NT2RP3000866	3.65	3.07	3.41	3.79	4.93	3.08	2.95	3.92	4.36			
30	NT2RP3000868	6.63	4.07	4.55	6.52	6.19	4.40	5.59	4.36	6.01			
	NT2RP3000869	7.38	5.89	6.47	6.37	7.71	6.66	5.72	5.36	5.4			
	NT2RP3000871	2.80	1.69	2.21	3.13	2.44	2.63	2.19	2.91	2.3			
	NT2RP3000875	6.14	2.07	3.11	2.15	2.68	3.67	3.92	2.74	3.62			
	NT2RP3000895	3.27	2.20	2.57	3.83	6.39	6.15	3.73	2.67	3.88	*	+	
	NT2RP3000900	9.85	5.60	5.12	11.99	12.50	10.94	7.71	7.19	8.22	*	+	
35	NT2RP3000901	5.01	2.45	2.11	6.45	8.36	6.11	4.49	5.69	7.42	*	+	
	NT2RP3000903	2.28	1.60	1.75	4.44	6.62	5.24	4.43	2.98	3.76	**	+	*
	NT2RP3000904	2.30	1.61	2.05	2.19	1.89	3.97	2.54	3.22	2.14			
	NT2RP3000907	9.61	6.08	7.44	8.62	11.64	8.56	8.91	8.78	9.69			
	NT2RP3000913	7.70	2.80	3.71	8.25	8.06	6.91	5.87	6.50	4.94			
40	NT2RP3000917	10.36	7.31	5.72	9.00	16.41	11.45	7.56	6.56	8.24			
	NT2RP3000919	5.76	4.04	3.02	5.13	7.71	4.25	4.75	6.45	6.91			
	NT2RP3000921	3.51	1.70	2.76	4.60	7.92	2.75	6.8	3.67	4.11			
	NT2RP3000942	9.61	5.52	5.34	12.62	14.38	12.46	6.8	6.53	7.24	*	+	
	NT2RP3000968	103.66	58.95	83.91	147.53	158.89	133.89	55.3	53.20	43.04	*	+	
45	NT2RP3000974	3.04	1.59	2.65	3.97	5.03	4.21	2.71	3.66	2.41	*	+	
	NT2RP3000980	39.62	20.55	29.98	6.47	9.37	6.00	4.91	6.99	8.46	*	-	*
	NT2RP3000984	5.29	4.18	5.73	10.16	10.11	7.87	6.25	8.85	4.44	**	+	
	NT2RP3000994	3.63	2.42	1.96	4.75	5.40	3.69	3.58	4.22	3.83			
	NT2RP3001001	3.47	2.25	3.10	3.83	2.41	2.13	2.68	3.98	2.58			
	NT2RP3001004	1.80	1.40	1.87	2.71	2.31	1.48	2.16	4.18	3			
50	NT2RP3001007	4.63	2.03	2.66	14.00	6.75	8.49	6.39	6.25	5.07	*	+	*
	NT2RP3001012	5.10	1.75	3.11	5.04	4.34	5.34	2.86	4.75	2.29			
	NT2RP3001042	5.71	3.43	4.72	5.27	4.96	3.88	3.98	3.86	2.98			
	NT2RP3001044	7.02	3.73	5.60	14.85	12.04	12.37	9.89	10.94	7.73	**	+	*
	NT2RP3001048	2.35	1.96	3.94	3.25	4.98	4.26	3.16	2.56	3.24			
55	NT2RP3001050	11.91	8.75	3.68	7.09	10.52	7.57	19.34	10.54	18.84			
	NT2RP3001055	19.61	12.87	10.53	9.87	9.64	7.47	11.2	7.71	10.89			

Table 268

5	NT2RP3000460	6.78	3.61	3.73	8.08	8.65	7.41	13.4	12.29	10.37	*	+	**	+
	NT2RP3000471	6.95	4.34	4.50	7.79	8.60	6.26	4.55	7.24	5.12				
	NT2RP3000477	21.65	12.36	-9.87	23.85	19.48	15.72	11.17	14.00	11.16				
	NT2RP3000478	7.29	4.54	5.34	13.47	15.94	11.73	8.07	4.43	7.92	**	+		
	NT2RP3000481	0.63	0.59	0.73	1.35	1.95	1.38	0.46	2.40	1.02	*	+		
	NT2RP3000484	1.55	0.72	1.25	1.68	2.10	2.87	1.12	2.90	1.09				
10	NT2RP3000487	5.07	1.99	2.06	3.79	5.91	4.35	2.41	2.16	2.61				
	NT2RP3000512	6.71	4.34	3.46	3.23	5.10	5.08	2.77	4.20	4.93				
	NT2RP3000523	27.58	15.65	17.30	17.42	22.63	15.01	11.77	10.31	9.03				
	NT2RP3000526	2.57	1.90	3.01	5.30	4.16	4.98	2.88	5.37	3.11	**	+		
	NT2RP3000527	3.80	1.53	2.25	4.05	4.14	5.85	2.46	3.30	2.31				
15	NT2RP3000531	15.89	10.13	8.97	23.60	23.41	21.43	13.33	15.19	15.55	**	+		
	NT2RP3000532	6.87	3.91	4.69	7.54	6.97	6.82	3.54	4.64	3.97				
	NT2RP3000542	4.26	2.58	3.40	6.33	6.95	7.50	5.58	5.25	4.09	**	+		
	NT2RP3000554	21.26	8.36	10.64	9.79	12.63	8.67	7.85	5.66	7.16				
	NT2RP3000561	1.72	1.29	0.49	4.36	4.39	2.75	5.41	6.15	4.61	*	+	**	+
20	NT2RP3000562	5.35	3.52	2.70	6.24	5.67	6.85	4.69	5.36	4.51	*	+		
	NT2RP3000578	2.48	1.13	0.91	1.41	2.33	1.20	1.51	2.72	1.83				
	NT2RP3000582	2.70	1.06	2.14	1.55	1.76	2.00	1.13	2.91	1.43				
	NT2RP3000584	3.87	1.71	2.00	3.83	3.38	4.15	1.95	3.50	3.43				
	NT2RP3000586	4.68	3.18	3.48	5.21	5.82	4.88	4.06	4.66	4.73	*	+		
25	NT2RP3000590	3.21	1.61	2.30	2.02	1.87	2.52	1.95	2.50	2.25				
	NT2RP3000592	2.67	1.26	1.45	1.25	2.76	1.46	1.33	1.90	1.13				
	NT2RP3000596	20.65	9.80	8.82	23.94	26.59	16.13	11.86	9.91	14.07				
	NT2RP3000599	3.31	1.41	2.33	3.96	4.14	2.63	2.43	4.34	3.3				
	NT2RP3000603	4.81	2.59	2.37	5.30	5.93	6.54	3.73	4.56	4.65	*	+		
	NT2RP3000605	2.51	1.85	1.50	3.30	3.59	2.96	2.17	4.09	3.29	*	+		
30	NT2RP3000607	7.51	5.55	8.79	5.67	5.09	3.67	3.76	3.78	3.57			*	-
	NT2RP3000616	2.94	0.94	1.60	3.25	4.41	3.35	2.18	3.01	2.34				
	NT2RP3000621	4.36	2.30	3.65	4.44	7.67	4.30	4.7	5.31	5.41				
	NT2RP3000622	6.01	4.28	3.80	5.09	7.11	5.45	5.08	3.73	3.94				
	NT2RP3000624	7.72	5.67	3.32	6.67	8.14	5.52	5.24	3.13	5.14				
35	NT2RP3000628	7.54	4.50	3.20	10.58	21.80	10.94	10.01	5.74	10.27				
	NT2RP3000631	16.09	7.17	9.25	14.57	17.16	15.18	7.31	8.71	8.97				
	NT2RP3000632	7.31	3.75	5.02	6.89	10.18	9.21	4.07	4.61	4.79				
	NT2RP3000638	7.68	5.11	4.32	4.07	4.85	4.59	5.9	6.86	5.24				
	NT2RP3000644	19.00	10.57	14.03	22.53	22.68	23.63	19.34	20.74	17.56	*	+		
40	NT2RP3000645	22.63	12.76	16.07	25.22	24.49	30.53	19.65	22.44	19.81				
	NT2RP3000652	25.30	13.23	15.28	45.18	43.44	33.63	16.17	15.59	14.77	*	+		
	NT2RP3000658	10.87	4.38	5.61	9.08	8.70	4.57	4.84	5.59	6.4				
	NT2RP3000660	7.86	3.20	4.43	11.71	10.96	7.67	5.63	5.73	5.08				
	NT2RP3000661	5.33	3.07	4.20	8.73	10.09	5.63	4.67	5.28	4.19				
45	NT2RP3000665	6.64	1.93	2.75	5.80	4.45	4.67	4.17	5.21	4.12				
	NT2RP3000676	8.20	4.06	3.78	8.46	10.33	8.20	6.83	8.27	6.88				
	NT2RP3000677	4.44	2.49	3.08	10.60	15.84	15.84	2.62	4.06	2.32	**	+		
	NT2RP3000681	16.25	8.48	11.24	17.10	13.94	12.61	11.39	15.24	10.7				
	NT2RP3000683	10.17	2.34	3.24	19.41	15.14	11.09	6.65	5.82	9.12	*	+		
	NT2RP3000685	7.81	3.42	2.68	6.13	4.88	5.09	4.14	3.91	7.49				
50	NT2RP3000690	3.45	1.81	2.38	2.69	3.42	3.19	1.6	4.35	4				
	NT2RP3000698	3.44	1.71	1.90	3.98	4.36	3.04	3.05	5.26	3.03				
	NT2RP3000708	8.35	3.44	2.85	6.09	5.53	5.09	2.92	4.17	5.63				
	NT2RP3000719	6.12	2.90	4.00	7.25	5.34	4.25	3.6	5.95	3.12				
	NT2RP3000721	4.08	2.25	2.01	4.97	4.56	3.47	2.13	3.08	2.89				
55	NT2RP3000728	2.25	0.64	0.87	2.34	2.75	2.13	0.67	2.18	0.8				
	NT2RP3000730	1.35	0.93	1.10	1.92	2.14	1.30	2.2	1.20	1.35				
	NT2RP3000733	4.35	2.50	1.71	6.01	6.36	4.79	3.49	3.48	2.85	*	+		

Table 267

5	NT2RP3000232	7.80	2.59	4.87	14.07	13.78	10.58	4.43	6.16	6.42	*	+		
	NT2RP3000233	4.29	2.04	3.30	4.16	4.02	3.58	3.88	4.05	3.95				
	NT2RP3000234	5.82	3.69	3.99	6.88	6.24	5.76	5.09	5.25	5.52				
	NT2RP3000235	4.07	2.16	2.75	4.46	3.39	3.79	3.35	4.56	3.36				
	NT2RP3000239	7.80	3.65	4.61	5.36	6.98	5.05	3.92	4.89	7.01				
	NT2RP3000247	2.30	1.21	1.95	2.12	2.01	2.94	1.85	3.86	2.35				
10	NT2RP3000251	8.89	5.54	6.24	11.87	10.35	8.87	9.19	9.33	7.77				
	NT2RP3000252	15.04	4.46	4.08	9.00	9.21	5.83	5.52	5.10	6.74				
	NT2RP3000255	5.13	2.85	2.23	3.53	3.93	3.32	2.12	2.52	4.46				
	NT2RP3000262	7.20	3.34	3.67	7.23	8.28	5.67	4.7	4.54	3.81				
	NT2RP3000266	13.99	6.47	5.93	16.36	16.88	13.38	9.91	14.29	13.15				
15	NT2RP3000267	4.19	1.73	1.51	3.08	4.17	2.39	2.28	3.19	2.31				
	NT2RP3000271	7.47	3.16	2.85	7.84	6.39	5.57	3.5	5.30	3.75				
	NT2RP3000278	3.14	2.04	3.02	4.79	6.42	5.26	3.22	5.70	4.21	**	+		
	NT2RP3000281	7.14	3.51	4.30	9.39	7.57	6.94	6.62	8.48	7.76				
	NT2RP3000292	2.43	1.31	1.46	1.66	2.08	1.80	2.97	2.36	1.82				
20	NT2RP3000299	3.32	1.72	2.64	3.50	2.85	1.65	3.49	2.65	2.85				
	NT2RP3000304	7.20	4.06	3.87	3.27	5.90	6.50	4.23	4.68	5.46				
	NT2RP3000310	9.88	5.44	4.97	10.57	8.79	8.65	8.38	7.53	9.91				
	NT2RP3000312	4.71	2.11	3.36	4.19	4.91	4.91	2.11	3.53	4.02				
	NT2RP3000320	9.82	2.79	5.46	8.18	6.79	9.80	7.95	7.10	16.94				
	NT2RP3000322	30.65	18.22	26.99	58.85	49.93	31.40	36.14	39.97	34.74		*	+	
25	NT2RP3000324	2.18	1.49	1.41	2.10	2.20	2.50	2.87	1.62	1.63				
	NT2RP3000326	4.07	2.09	2.65	6.40	4.79	6.20	5.05	3.50	3.68	*	+		
	NT2RP3000329	8.08	3.03	2.39	13.04	10.42	8.93	5.43	5.08	6.48	*	+		
	NT2RP3000330	6.13	3.81	4.47	3.99	4.93	3.61	5.76	6.52	5.37				
	NT2RP3000333	3.58	1.99	1.19	2.09	2.88	2.04	2.14	2.57	2.31				
30	NT2RP3000341	13.34	6.74	7.40	16.98	14.13	16.48	11.16	11.51	12.58	*	+		
	NT2RP3000344	2.19	2.15	1.77	2.27	1.91	1.50	1.56	1.76	2.32				
	NT2RP3000345	0.88	0.64	0.51	3.07	2.22	3.27	0.95	0.77	2.11	**	+		
	NT2RP3000348	112.18	53.12	48.19	87.36	67.82	76.37	170.4	141.05	175.2		*	+	
	NT2RP3000350	13.69	7.30	6.99	9.25	9.00	7.77	7.42	5.74	8.01				
35	NT2RP3000359	10.64	6.49	5.35	19.00	17.38	16.68	15.5	13.49	16.08	**	+	*	+
	NT2RP3000361	10.35	4.92	4.34	11.24	6.97	7.55	6.16	6.69	7.28				
	NT2RP3000366	7.65	3.30	4.82	9.45	14.23	10.18	10.84	11.42	12.66	*	+	*	+
	NT2RP3000378	4.91	3.67	4.88	5.34	6.49	6.00	4.34	4.99	3.64				
	NT2RP3000384	6.56	5.43	5.50	8.93	9.13	11.76	6.91	6.90	7.16	*	+	*	+
	NT2RP3000389	14.26	10.15	11.05	22.04	27.40	18.38	12.47	13.44	23.39	*	+		
40	NT2RP3000393	5.27	3.15	2.77	4.98	4.37	4.43	4.32	3.00	3.71				
	NT2RP3000395	121.26	84.54	65.25	98.14	119.90	103.24	32.56	26.84	40.17		*	-	
	NT2RP3000397	3.69	4.24	2.44	2.76	4.13	3.97	3.48	2.62	4.13				
	NT2RP3000398	6.97	4.09	4.94	8.35	10.97	6.66	5.51	6.21	5.86				
	NT2RP3000403	4.82	3.83	4.35	9.87	12.59	8.19	6.65	6.56	8.79	*	+	*	+
45	NT2RP3000418	4.00	2.62	2.61	8.58	12.65	8.62	5.36	6.28	7.5	**	+	*	+
	NT2RP3000424	5.08	4.11	3.96	14.10	16.88	10.90	8.47	7.77	7.95	**	+	**	+
	NT2RP3000427	2.50	1.80	2.77	5.73	6.63	8.27	3.99	5.02	3.87	**	+	*	+
	NT2RP3000431	3.51	2.32	1.35	4.97	4.03	2.77	4.39	4.52	3.47				
	NT2RP3000433	4.48	3.35	3.32	4.96	5.89	5.97	3.9	4.05	4.56	*	+		
50	NT2RP3000436	11.10	6.79	5.78	9.34	10.99	9.24	10.36	9.52	16.87				
	NT2RP3000439	5.21	2.28	3.00	3.90	7.56	3.69	3.69	4.00	3.42				
	NT2RP3000441	1.19	0.92	0.83	1.64	2.07	1.50	2.8	3.37	2.81	*	+	**	+
	NT2RP3000444	2.26	2.00	1.85	2.13	2.91	3.48	2.82	2.26	2.53				
	NT2RP3000448	3.48	2.24	3.61	8.12	11.89	8.40	5.13	4.03	6.51	**	+		
	NT2RP3000449	5.49	2.45	3.20	2.67	4.04	3.28	1.61	2.66	1.96				
55	NT2RP3000451	5.47	3.68	2.74	2.86	3.50	4.17	4.01	4.24	4.31				
	NT2RP3000456	4.82	4.21	3.70	3.94	5.59	4.96	4.41	3.70	5.42				

Table 266

5	NT2RP2006554	2.93	1.44	1.64	4.14	5.11	5.65	3.05	2.87	4.34	**	+		
	NT2RP2006565	2.42	3.04	1.97	5.84	7.27	4.73	5.76	4.50	8.32	*	+	*	+
	NT2RP2006571	15.53	8.80	8.87	9.19	10.25	5.31	9.49	9.09	15.1				
	NT2RP2006573	3.03	1.23	1.11	3.74	3.96	3.02	2.6	2.13	2.11				
	NT2RP2006598	5.73	3.98	4.61	7.93	8.72	6.43	5.28	3.71	6.12	*	+		
	NT2RP2006601	37.52	34.93	32.64	41.04	41.47	32.68	27.39	28.66	36.43				
10	NT2RP3000002	3.95	2.25	3.29	4.37	7.61	7.60	3.47	4.83	7.96	*	+		
	NT2RP3000011	4.07	2.64	1.62	5.92	4.70	5.14	3.96	3.12	4.24	*	+		
	NT2RP3000014	3.17	3.00	2.39	9.14	11.05	8.39	7.15	7.48	8.57	**	+	**	+
	NT2RP3000016	9.66	5.49	5.68	6.73	6.36	7.49	4.75	5.66	6.35				
	NT2RP3000022	4.96	2.03	2.47	3.53	3.43	2.45	3.24	3.89	7.1				
15	NT2RP3000024	12.74	9.32	13.69	28.77	37.69	22.23	11.49	12.80	14.79	*	+		
	NT2RP3000031	4.64	2.28	2.98	4.90	4.09	5.50	4.12	3.94	3.26				
	NT2RP3000034	4.51	3.69	3.49	3.95	4.58	4.75	3.38	3.05	3.23				
	NT2RP3000037	15.49	9.32	10.69	13.56	14.15	12.81	7.78	9.45	8.16				
	NT2RP3000040	2.98	2.45	1.73	1.43	1.95	2.12	0.99	2.09	2.1				
20	NT2RP3000041	10.75	6.47	4.78	19.57	16.79	13.38	9.67	7.12	9.17	*	+		
	NT2RP3000046	5.16	2.85	2.89	6.40	9.13	5.39	4.23	3.75	6.16				
	NT2RP3000047	6.44	3.75	3.07	4.50	4.32	4.37	3.44	4.24	4.69				
	NT2RP3000049	3.94	3.36	1.85	3.67	6.35	6.22	5.02	4.43	8.2				
	NT2RP3000050	7.94	4.67	6.52	13.03	15.60	12.76	7.92	7.66	10.86	**	+		
25	NT2RP3000051	6.26	3.23	4.99	9.29	9.59	8.78	5.46	7.17	6.65	**	+		
	NT2RP3000054	6.09	3.47	4.38	5.67	6.99	5.26	5.01	4.84	5.62				
	NT2RP3000055	3.24	2.73	0.81	4.89	4.66	2.53	2.67	2.43	3.79				
	NT2RP3000056	2.70	3.24	1.60	2.60	3.66	2.74	3.75	2.94	3.3				
	NT2RP3000059	4.21	2.87	2.12	3.45	3.50	3.02	3.35	3.22	4.21				
	NT2RP3000063	7.78	5.44	6.74	6.64	5.14	7.47	6.5	8.34	4.12				
30	NT2RP3000068	1.30	1.86	2.21	1.64	3.20	2.26	2.1	3.07	3.12				
	NT2RP3000069	3.21	2.16	2.26	10.79	10.68	7.75	8.64	7.90	7.98	**	+	**	+
	NT2RP3000072	2.08	1.15	1.36	3.34	2.75	2.73	2.05	3.07	2.12	*	+		
	NT2RP3000080	12.90	8.84	11.62	14.83	16.14	12.41	14.4	11.56	12.15				
	NT2RP3000085	4.82	2.44	2.00	2.73	3.07	3.01	2.95	2.26	2.49				
35	NT2RP3000087	12.35	7.36	5.97	19.26	20.25	18.12	12.89	8.99	10.11	**	+		
	NT2RP3000092	2.83	2.11	1.59	4.04	2.45	1.56	2.71	2.87	2.87				
	NT2RP3000109	1.75	1.89	2.71	5.02	4.39	3.90	1.58	3.14	1.97	**	+		
	NT2RP3000119	10.48	4.74	6.30	7.48	8.15	6.85	5.44	7.67	7.52				
	NT2RP3000125	9.53	6.24	6.75	10.54	13.59	12.33	7.17	8.82	6.83	*	+		
40	NT2RP3000131	13.37	7.84	8.67	12.43	13.75	13.12	11.27	10.91	10.26				
	NT2RP3000134	8.39	4.00	4.04	11.86	8.47	11.09	6.57	5.88	5.18	*	+		
	NT2RP3000137	7.33	3.86	4.11	4.55	10.23	5.70	5.38	4.77	5.25				
	NT2RP3000142	8.58	2.85	3.30	8.25	6.01	4.98	4.9	4.68	4.51				
	NT2RP3000148	6.50	3.03	2.82	4.77	5.93	4.35	4.39	4.87	3.31				
	NT2RP3000149	7.40	4.34	3.38	4.95	6.06	4.71	3.65	4.88	5.43				
45	NT2RP3000163	5.34	2.10	2.73	5.49	7.84	4.53	2.61	3.70	2.68				
	NT2RP3000168	17.73	9.34	8.35	13.43	12.52	14.26	15.5	18.99	23.37				
	NT2RP3000169	2.79	1.47	1.93	3.28	2.66	2.80	2.69	4.02	3.92				
	NT2RP3000171	30.99	20.17	24.95	41.61	37.53	33.55	22.47	25.44	33.88	*	+		
	NT2RP3000172	5.29	2.13	2.18	3.70	4.85	1.88	2.31	1.91	2.23				
50	NT2RP3000186	16.37	8.43	6.94	11.35	12.10	6.88	5.69	5.57	6.97				
	NT2RP3000197	2.96	2.49	2.66	5.21	6.67	3.78	2.54	2.96	3.64	*	+		
	NT2RP3000201	11.54	5.67	6.73	11.59	11.99	10.04	5.11	5.52	10.33				
	NT2RP3000204	3.53	2.05	1.72	2.68	3.65	3.34	1.75	3.41	1.98				
	NT2RP3000207	4.88	2.36	2.46	3.16	3.56	3.29	4.1	5.13	6.04				
55	NT2RP3000216	8.62	6.38	5.44	6.42	8.59	6.54	8.63	6.16	7.72				
	NT2RP3000220	2.88	1.23	2.50	2.75	3.70	2.71	2.57	3.23	2.33				
	NT2RP3000221	4.47	2.97	2.52	4.75	5.82	4.37	3.89	3.79	4.34				

Table 265

	NT2RP2006141	5.76	3.11	3.17	3.84	5.50	4.54	3.67	3.75	3.95				
5	NT2RP2006166	7.93	5.66	5.17	12.63	13.99	9.56	6.76	6.08	6.36	*	+		
	NT2RP2006176	4.45	2.26	-1.67	6.40	4.88	5.22	2.44	3.34	5.68	*	+		
	NT2RP2006181	1.58	1.06	1.00	1.37	3.24	3.22	1.23	2.94	1.73				
	NT2RP2006184	23.94	15.54	16.09	22.96	21.00	23.09	17.11	19.55	14.56				
	NT2RP2006186	1.68	1.14	2.35	2.02	3.74	1.74	1.23	3.31	1.82				
10	NT2RP2006196	4.74	3.02	3.70	6.83	6.02	5.77	4.04	5.17	3.91	*	+		
	NT2RP2006199	2.29	2.59	2.52	3.33	3.50	4.30	2.88	2.76	2.12	*	+		
	NT2RP2006200	4.29	2.63	1.43	3.59	5.59	2.06	3.12	2.50	2.5				
	NT2RP2006210	59.40	41.07	36.68	33.45	39.27	22.58	11.72	8.31	9.47		**		
	NT2RP2006219	3.75	1.76	1.64	3.39	3.29	2.82	2.17	1.88	4.22				
15	NT2RP2006224	5.72	3.72	4.01	5.11	6.26	6.39	3.82	3.77	4.2				
	NT2RP2006237	5.09	3.91	5.00	9.00	7.92	10.28	5.01	5.76	5.27	**	+		
	NT2RP2006238	3.42	2.16	1.78	4.42	4.29	2.44	2.31	3.01	1.89				
	NT2RP2006258	9.12	5.55	6.28	6.08	7.62	7.68	6.35	6.07	3.93				
	NT2RP2006261	1.75	2.42	1.14	2.06	2.49	1.87	1.21	1.75	2.67				
20	NT2RP2006269	23.86	9.30	9.53	15.39	18.13	13.53	12.46	10.61	15.67				
	NT2RP2006275	4.68	2.71	2.12	3.99	3.20	2.45	2.89	2.81	3.34				
	NT2RP2006282	7.12	3.89	6.34	8.17	11.45	9.25	4.48	4.87	2.85	*	+		
	NT2RP2006302	4.86	2.69	3.31	10.51	10.47	9.06	10	9.83	3.76	**	+		
	NT2RP2006312	8.45	5.62	5.99	10.60	10.03	9.84	7.18	6.51	5.02	*	+		
	NT2RP2006320	3.62	2.45	1.39	4.62	5.47	5.86	2.21	4.05	3.23	*	+		
25	NT2RP2006321	1.99	1.78	2.42	3.22	4.24	2.52	1.97	3.17	2.07				
	NT2RP2006323	1.30	0.75	0.38	1.35	1.65	0.69	0.19	2.09	2.6				
	NT2RP2006333	2.18	0.70	0.66	2.51	1.88	1.17	1.49	1.76	2.35				
	NT2RP2006334	3.73	1.40	1.47	2.69	3.03	2.34	0.81	2.29	2.95				
	NT2RP2006338	2.65	1.82	1.03	3.45	4.02	2.81	1.6	3.69	2.93				
30	NT2RP2006339	2.37	1.54	1.37	3.09	2.39	1.47	1.2	2.47	2.21				
	NT2RP2006355	1.01	0.99	0.71	2.16	2.25	1.72	1.94	2.95	0.87	**	+		
	NT2RP2006365	1.51	1.66	1.15	3.16	4.39	3.70	1.83	4.13	1.9	**	+		
	NT2RP2006374	16.70	8.19	7.22	17.36	18.00	12.60	10.86	13.62	9.02				
35	NT2RP2006393	4.85	2.17	2.52	8.54	10.40	8.85	5.98	6.32	6.15	**	+	*	+
	NT2RP2006394	2.02	1.64	1.69	3.46	1.86	1.52	3.53	1.56	2.54				
	NT2RP2006400	1.99	1.74	1.43	2.29	2.67	2.67	2.79	1.32	1.33	*	+		
	NT2RP2006411	36.13	23.40	20.23	18.85	35.68	22.21	26.26	22.92	21.44				
	NT2RP2006429	3.49	1.96	1.56	18.22	22.80	21.81	7.61	6.72	8.71	**	+	**	+
	NT2RP2006435	2.88	2.61	2.07	4.19	4.16	3.86	3.51	3.02	3.91	**	+		
40	NT2RP2006436	4.50	2.57	2.37	11.47	10.14	10.45	15.06	14.83	12.68	**	+	**	+
	NT2RP2006441	5.48	3.11	4.37	12.23	11.44	10.95	9.38	9.44	9.01	**	+	**	+
	NT2RP2006447	3.63	2.74	2.87	7.53	5.11	1.37	2.09	1.56	0.94		*		-
	NT2RP2006454	3.45	1.48	1.32	2.04	2.21	2.24	3.02	1.84	0.51				
	NT2RP2006455	3.08	1.02	1.42	3.46	1.52	2.11	2.25	1.46	1.25				
	NT2RP2006456	3.43	1.56	1.38	1.87	3.29	2.20	1.39	3.00	3.52				
45	NT2RP2006464	7.78	4.38	3.90	5.55	4.82	4.88	3.6	3.54	5.67				
	NT2RP2006467	5.66	2.72	2.67	10.90	8.83	10.03	7.29	6.50	9.64	**	+	*	+
	NT2RP2006472	7.44	3.78	3.97	8.69	8.19	8.22	5	4.65	10.62				
	NT2RP2006474	8.86	5.98	7.97	27.71	30.65	24.91	30	37.03	33.44	**	+	**	+
	NT2RP2006475	5.74	3.11	2.17	15.80	11.04	13.89	8.72	6.46	9.93	**	+	*	+
50	NT2RP2006476	14.81	5.32	5.83	6.07	6.15	5.06	6.6	4.52	6.65				
	NT2RP2006501	10.57	4.49	3.64	10.98	10.15	9.25	4.35	4.19	5.76				
	NT2RP2006512	10.18	4.42	5.26	7.98	9.45	6.77	5.81	5.75	6.47				
	NT2RP2006526	2.38	0.63	1.13	1.33	3.17	1.44	1.31	2.50	2.57				
	NT2RP2006527	6.04	4.50	5.90	6.98	6.77	8.30	6.37	6.06	6.05				
	NT2RP2006534	1.08	0.58	0.52	1.10	1.90	2.81	1.51	1.54	1.55		*		+
55	NT2RP2006537	7.96	4.17	4.11	12.78	11.80	12.98	5.84	7.01	9.26	**	+		
	NT2RP2006543	2.53	2.49	1.25	3.82	3.98	2.63	5.74	3.55	4.9		*		+

Table 264

5	NT2RP2005775	4.39	1.98	1.42	2.12	2.56	2.56	2.19	2.08	1.67				
	NT2RP2005781	5.85	3.98	3.29	6.76	5.57	5.04	4.75	3.50	4.17				
	NT2RP2005784	11.14	6.73	-5.29	8.15	8.38	8.40	7.85	8.40	10.24				
	NT2RP2005789	4.85	3.33	3.28	5.63	7.04	4.46	3.88	3.70	4.09				
	NT2RP2005799	1.71	1.81	1.37	3.76	5.36	2.16	2.16	2.19	2.43		*	+	
	NT2RP2005804	6.19	3.18	3.30	4.57	7.49	6.42	5.55	5.88	4.72				
10	NT2RP2005812	3.92	3.04	2.54	4.78	6.17	3.21	2.98	4.18	4.04				
	NT2RP2005815	2.54	2.17	3.20	3.81	3.69	2.58	2.35	2.98	1.88				
	NT2RP2005835	14.04	7.44	6.79	14.50	10.00	10.84	9.86	7.11	11.61				
	NT2RP2005841	6.35	3.23	3.13	5.70	4.93	4.82	5.84	3.68	4.27				
	NT2RP2005853	3.23	3.29	2.96	6.28	6.53	5.74	4.87	4.09	5.28	**	+	*	+
15	NT2RP2005857	8.95	4.28	4.74	6.65	7.52	6.19	1.63	2.12	1.8				
	NT2RP2005859	5.38	4.41	5.54	4.28	5.42	3.86	2.87	3.84	3.87		*	-	
	NT2RP2005860	3.02	1.60	2.64	2.92	4.01	2.37	2.32	4.74	1.81				
	NT2RP2005863	4.66	2.88	2.88	3.96	3.85	3.93	2.02	2.05	1.69				
	NT2RP2005868	3.44	1.57	1.65	4.52	4.28	2.97	2.38	3.85	2.89				
20	NT2RP2005876	13.61	7.01	5.40	17.03	13.16	6.91	8.8	8.61	107				
	NT2RP2005878	6.92	4.37	4.13	11.06	12.33	11.73	5.81	7.81	6.82	**	+		
	NT2RP2005883	1.59	1.56	1.08	3.31	2.84	2.42	3.91	4.53	4.86	**	+	**	+
	NT2RP2005886	8.60	4.98	6.40	10.11	11.16	11.42	6.19	6.08	5.43	*	+		
	NT2RP2005887	5.47	3.26	3.97	12.05	12.81	9.32	13.75	10.35	15.02	**	+	**	+
	NT2RP2005890	7.74	6.08	7.50	6.23	6.35	4.71	2.57	2.56	1.86		**	-	
25	NT2RP2005901	3.39	2.76	2.57	3.81	4.07	4.20	2.43	3.04	3.13	*	+		
	NT2RP2005902	1.86	0.89	1.33	3.39	3.77	2.15	2.13	2.79	3.13	*	+	*	+
	NT2RP2005908	9.46	5.71	4.03	9.28	7.93	10.45	6.03	6.26	6.92				
	NT2RP2005927	7.43	5.84	5.10	9.51	9.65	7.14	3.72	5.75	4.41				
	NT2RP2005933	6.32	4.20	3.63	5.57	7.02	4.50	3.29	2.73	4.08				
30	NT2RP2005941	9.03	6.94	7.01	7.65	13.07	8.78	10.41	9.47	5.87				
	NT2RP2005942	3.02	2.03	1.79	3.90	4.09	3.96	2.56	2.68	2	*	+		
	NT2RP2005946	6.57	4.95	5.93	3.90	3.86	3.27	2.5	2.94	2.41	*	-	**	-
	NT2RP2005970	12.30	10.25	11.94	15.87	16.05	15.06	14.9	13.37	14.97	**	+	*	+
	NT2RP2005980	3.71	2.65	2.25	7.90	7.37	4.49	4.13	4.23	2.71	*	+		
35	NT2RP2005994	5.01	2.60	2.01	2.75	4.22	1.07	2.23	3.11	2.43				
	NT2RP2006004	2.32	1.82	1.35	2.43	4.21	2.56	2.36	3.37	2.03				
	NT2RP2006013	4.44	2.15	4.45	6.09	6.99	3.28	4.68	5.22	4.41				
	NT2RP2006023	21.60	12.40	20.04	37.44	49.33	45.44	22.61	22.79	24.39	**	+		
	NT2RP2006028	5.34	3.20	3.73	4.07	4.23	2.81	3.39	4.81	5.42				
40	NT2RP2006038	0.34	0.06	1.28	0.43	0.83	3.61	0.25	1.80	0.18				
	NT2RP2006042	8.65	5.14	6.93	7.32	7.79	6.34	7.56	7.82	9.4				
	NT2RP2006043	5.05	2.75	2.80	12.32	12.87	10.73	8.05	8.08	7.82	**	+	**	+
	NT2RP2006052	2.31	2.64	1.44	1.42	2.55	2.98	1.26	2.10	2.6				
	NT2RP2006057	3.69	1.67	1.24	3.57	3.44	2.48	2.2	3.85	3.27				
	NT2RP2006064	12.49	6.77	9.83	12.13	10.85	6.00	10.28	6.81	5.57				
45	NT2RP2006068	3.25	3.63	2.31	8.60	6.86	6.64	4.6	5.09	2.54	**	+		
	NT2RP2006069	1.08	0.69	0.92	0.88	1.74	0.95	0.92	1.48	1.42				
	NT2RP2006071	2.73	3.23	2.31	5.07	7.66	5.45	2.92	4.00	2.55	*	+		
	NT2RP2006090	3.70	1.69	2.79	3.57	5.20	3.82	3.74	3.63	2.49				
	NT2RP2006092	3.65	2.47	2.47	3.19	3.41	3.44	2.36	2.80	2.77				
50	NT2RP2006097	24.23	9.76	10.66	21.53	18.65	14.12	10.2	9.65	12.96				
	NT2RP2006098	4.17	2.27	1.77	4.26	4.04	2.86	4.03	6.51	3.5				
	NT2RP2006099	4.48	2.99	2.12	5.82	5.86	5.18	3.32	4.84	3.86	*	+		
	NT2RP2006100	3.88	1.55	1.83	3.98	4.92	3.85	2.1	3.68	2.65				
	NT2RP2006183	10.54	3.86	5.78	3.37	2.55	1.88	2.21	3.36	1.85				
55	NT2RP2006106	8.45	4.11	4.04	6.45	5.69	6.30	4.48	6.29	4.22				
	NT2RP2006127	9.00	6.34	7.56	9.10	8.66	7.24	9.22	10.10	8.49				
	NT2RP2006134	1.55	1.02	1.47	1.76	1.82	1.93	1.55	2.52	1.29	*	+		

Table 263

	NT2RP2005498	6.78	2.60	2.45	2.62	6.63	3.50	3.33	3.34	4.18				
	NT2RP2005501	4.44	2.53	2.65	2.38	4.12	2.69	2.07	3.28	2.78				
5	NT2RP2005506	5.72	4.30	3.10	5.43	9.55	6.10	24.52	21.82	25.02		**	+	
	NT2RP2005509	6.91	5.58	4.63	12.32	11.78	9.14	5.34	8.99	8.48	*	+		
	NT2RP2005514	3.36	2.23	2.33	3.96	5.18	4.19	3.03	4.16	4.55	*	+		
	NT2RP2005520	10.34	5.10	5.86	6.07	8.22	5.46	3.87	3.79	3.08				
	NT2RP2005525	6.12	4.01	5.33	8.58	7.75	8.13	5.26	8.01	5.47	*	+		
10	NT2RP2005531	0.65	1.10	1.57	2.33	1.56	1.74	1.49	2.39	1.21				
	NT2RP2005535	36.57	17.31	21.13	93.90	73.03	67.87	27.53	17.14	25.99	**	+		
	NT2RP2005539	10.87	6.53	4.81	8.43	9.17	6.85	6.76	6.87	5.25				
	NT2RP2005540	2.81	2.63	2.81	7.15	6.27	5.67	4.42	5.46	9.74	**	+		
	NT2RP2005541	5.40	3.42	2.70	8.82	9.81	10.04	7.49	7.37	5.44	**	+	*	+
15	NT2RP2005549	3.91	1.98	1.81	3.23	3.51	2.41	2.43	3.46	2.97				
	NT2RP2005555	3.52	2.33	3.66	6.38	7.55	5.49	7.54	10.56	6.47	*	+	*	+
	NT2RP2005557	7.00	5.12	11.72	16.35	11.47	12.41	6.34	5.80	8.04				
	NT2RP2005581	5.51	4.09	4.45	13.70	13.23	10.54	6.26	5.62	5.86	**	+		
	NT2RP2005586	7.40	3.49	4.35	2.55	4.08	2.63	1.67	2.60	2.43				
20	NT2RP2005597	6.16	4.97	3.02	4.57	4.34	4.57	4.67	4.40	5.08				
	NT2RP2005600	4.06	2.52	2.53	3.83	4.26	3.10	2.47	4.00	2.95				
	NT2RP2005605	13.12	8.01	6.74	12.67	14.30	12.26	6.96	7.51	8				
	NT2RP2005614	9.18	5.27	8.25	16.39	16.00	13.57	10.11	8.70	9.2	**	+		
	NT2RP2005620	4.07	2.65	2.40	3.99	3.40	3.40	2.45	3.61	2.26				
25	NT2RP2005622	9.20	6.36	7.23	6.07	7.94	5.76	4.64	4.67	6.34				
	NT2RP2005632	3.64	3.42	2.57	5.77	4.33	3.82	2.82	3.85	3.3				
	NT2RP2005635	3.95	2.73	2.06	3.40	4.38	2.94	2.4	2.42	3.18				
	NT2RP2005637	2.20	1.05	1.68	13.21	4.02	4.55	2.2	2.55	5.6				
	NT2RP2005640	3.47	1.55	1.53	2.16	1.23	2.22	1.96	2.66	2.84				
	NT2RP2005645	6.42	3.67	2.99	5.68	11.68	7.34	5.29	6.74	5.73				
30	NT2RP2005651	4.09	3.02	3.19	6.89	11.77	5.52	3.81	4.33	6.7				
	NT2RP2005654	5.50	3.61	4.20	6.10	7.84	5.96	4.19	5.64	4.96				
	NT2RP2005666	4.54	3.08	3.45	5.18	6.63	4.14	4.25	3.69	7.2				
	NT2RP2005669	6.09	5.35	5.64	8.34	9.73	9.01	4.66	6.00	6.82	**	+		
	NT2RP2005670	2.87	2.37	1.87	5.75	5.68	2.37	1.68	2.33	3.03				
35	NT2RP2005671	10.41	3.42	4.33	5.10	6.32	3.51	3.46	4.47	6.12				
	NT2RP2005675	11.31	4.30	4.30	8.54	8.22	4.79	7.64	6.94	9.43				
	NT2RP2005683	9.32	5.43	5.87	8.08	9.48	5.92	5.85	4.94	4.56				
	NT2RP2005690	3.18	1.30	1.52	3.24	4.46	3.75	2.33	3.71	3.54				
	NT2RP2005694	4.33	2.30	2.18	4.82	3.54	4.62	3.22	3.77	3.78				
40	NT2RP2005701	22.21	13.84	17.86	22.12	25.56	24.08	18.18	17.70	22.41				
	NT2RP2005712	2.84	3.06	3.02	3.90	3.94	3.10	1.15	2.49	1.88		*	-	
	NT2RP2005719	2.26	1.27	0.73	3.09	3.04	2.67	2.23	1.46	2.56	*	+		
	NT2RP2005722	11.76	8.52	5.52	18.21	24.59	18.10	8.26	9.21	12.37	*	+		
	NT2RP2005723	4.68	2.75	2.29	7.35	6.52	3.86	4.39	4.70	2.79				
45	NT2RP2005726	5.41	2.39	2.73	5.77	4.51	4.16	3.27	4.19	3.67				
	NT2RP2005729	5.30	2.58	2.08	6.82	6.27	4.01	3.21	5.54	3.89				
	NT2RP2005731	0.50	0.60	0.63	1.06	1.43	0.80	0.71	2.81	0.87	*	+		
	NT2RP2005732	8.98	3.61	4.01	6.71	6.46	5.79	4.23	7.06	7.16				
	NT2RP2005737	10.83	8.16	10.12	14.65	17.80	12.60	12.9	11.51	9.06	*	+		
50	NT2RP2005741	5.83	2.63	2.65	3.36	3.80	2.41	3.96	2.72	3.47				
	NT2RP2005748	3.52	1.63	2.33	2.18	2.64	1.48	3.11	2.62	2.38				
	NT2RP2005752	5.37	3.43	3.73	6.46	5.65	5.66	6.55	3.67	3.82				
	NT2RP2005753	22.04	14.07	18.05	15.96	24.14	20.59	21.63	18.25	19.82				
	NT2RP2005763	6.73	2.47	2.52	3.25	3.61	3.70	1.84	3.88	3.22				
	NT2RP2005767	2.43	2.60	2.16	6.91	6.56	7.20	3.36	3.03	4.12	**	+	*	+
55	NT2RP2005773	15.62	10.12	12.99	19.66	19.02	17.26	17.15	13.07	15.8	*	+		
	NT2RP2005774	10.33	5.72	6.91	21.21	24.60	21.03	9.42	7.55	8.22	**	+		

Table 262

	NT2RP2005139	3.84	1.72	1.31	3.14	3.97	2.27	2.16	2.35	2.71			
	NT2RP2005140	6.44	3.34	1.76	2.06	2.19	1.94	1.62	2.45	4.48			
5	NT2RP2005144	7.59	4.23	3.57	8.56	9.25	7.68	4.75	8.24	8.15			
	NT2RP2005147	3.33	1.34	1.33	2.20	2.64	3.04	4.92	2.37	1.84			
	NT2RP2005148	4.87	2.83	2.05	4.55	5.06	4.19	2.73	4.23	3.35			
	NT2RP2005159	3.35	2.32	2.38	3.01	3.13	3.18	2.03	3.88	1.9			
	NT2RP2005162	3.09	1.68	1.72	3.70	3.44	2.30	2.24	3.35	2.16			
10	NT2RP2005163	25.94	15.25	17.25	21.49	24.77	28.25	17.62	25.86	21.18			
	NT2RP2005168	4.54	2.65	2.28	2.25	4.03	2.91	2.1	1.69	2.5			
	NT2RP2005181	9.05	4.31	4.53	4.26	4.18	3.03	3.8	2.76	3.1			
	NT2RP2005204	8.22	7.14	6.39	7.26	7.87	6.45	7	4.58	3.93			
	NT2RP2005219	6.43	4.48	4.74	6.61	6.15	4.27	4.15	5.58	7.21			
15	NT2RP2005227	6.13	3.78	3.14	9.09	11.14	7.97	3.82	5.07	8.88	*	+	
	NT2RP2005237	27.33	18.84	15.64	23.79	22.48	23.44	22.52	21.69	18.11			
	NT2RP2005239	3.74	1.34	1.71	2.73	2.86	2.63	2.66	2.69	2.3			
	NT2RP2005247	2.49	2.14	1.98	4.28	4.68	4.69	2.63	2.43	2.5	**	+	
	NT2RP2005254	9.04	3.29	3.29	8.47	7.53	8.80	7.01	6.79	4.08			
20	NT2RP2005270	4.99	2.71	2.82	6.57	6.85	4.80	6.2	6.16	8.3		*	+
	NT2RP2005276	9.47	6.54	6.31	10.41	11.77	12.24	5.39	7.57	7.48	*	+	
	NT2RP2005287	4.80	3.96	2.36	5.91	7.62	8.20	5.51	5.27	7.29	*	+	
	NT2RP2005288	3.78	1.10	1.91	4.67	4.69	3.22	2.56	2.68	2.46			
	NT2RP2005289	3.95	2.82	3.63	10.36	10.31	13.45	7.04	9.38	8.68	**	+	**
25	NT2RP2005293	4.69	3.98	2.48	2.80	6.37	4.36	1.98	2.19	8.18			
	NT2RP2005315	4.50	2.51	3.53	6.84	5.84	6.72	4.55	3.38	3.33	*	+	
	NT2RP2005322	8.85	3.21	3.77	5.49	9.42	5.85	5.53	11.41	21.87			
	NT2RP2005325	13.28	7.03	7.32	9.81	8.97	5.93	11.14	10.62	11.49			
	NT2RP2005336	12.73	6.78	5.54	13.58	10.27	12.67	8.85	6.83	5.91			
30	NT2RP2005343	6.02	1.89	2.05	7.45	9.65	7.01	10.08	10.85	12.82	*	+	**
	NT2RP2005344	1.85	1.66	1.47	2.08	2.88	1.92	2.74	2.45	3.15		**	+
	NT2RP2005347	4.37	2.71	1.89	5.25	5.00	4.78	3.35	2.93	2.34			
	NT2RP2005354	12.00	6.61	6.14	17.43	12.77	12.49	8.48	9.88	9.01			
	NT2RP2005358	4.88	3.45	2.64	4.51	4.14	3.14	3.97	2.53	1.99			
35	NT2RP2005360	7.88	5.76	2.39	6.48	5.68	6.59	4.31	3.84	6.35			
	NT2RP2005378	18.33	8.81	8.98	11.83	10.64	10.23	12.69	11.85	15.35			
	NT2RP2005391	11.21	5.99	4.87	8.42	9.50	6.15	7.72	6.42	7.6			
	NT2RP2005393	7.14	5.04	4.09	7.19	7.55	7.32	5.14	5.24	6.8			
	NT2RP2005407	4.70	3.27	2.59	4.12	5.86	4.29	4.19	4.07	6.46			
	NT2RP2005419	2.03	2.94	2.38	2.87	3.30	2.26	2.46	2.93	2.38			
40	NT2RP2005425	3.16	1.77	1.43	6.79	4.57	5.63	3.84	5.07	4.35	*	+	*
	NT2RP2005429	5.40	3.41	3.71	7.74	6.15	6.01	3.54	4.45	2.89	*	+	
	NT2RP2005436	11.49	5.63	5.95	16.34	13.38	12.70	9.59	8.94	10.22	*	+	
	NT2RP2005441	2.64	2.24	1.49	4.39	3.02	4.62	2.37	2.61	2.65	*	+	
	NT2RP2005442	6.72	3.80	3.11	6.35	6.53	5.07	6.08	5.33	7.07			
45	NT2RP2005444	14.62	10.40	7.75	7.08	9.17	7.55	7.37	6.88	8.45			
	NT2RP2005453	1.54	2.20	1.49	7.95	9.47	8.01	8.67	8.44	9.15	**	+	**
	NT2RP2005457	15.76	12.87	16.87	26.94	13.90	21.92	12.51	12.21	12.15			
	NT2RP2005458	1.63	1.87	2.03	5.92	5.93	3.89	2.67	3.64	6.17	**	+	
	NT2RP2005463	4.65	3.64	4.43	7.72	7.84	5.33	6.02	6.93	5.84	*	+	*
50	NT2RP2005464	11.98	9.14	6.68	11.62	10.20	8.75	5.59	3.86	4.74		*	
	NT2RP2005465	4.57	3.64	2.60	8.98	7.23	8.68	2.44	5.04	5.3	**	+	
	NT2RP2005472	10.01	4.28	4.30	7.95	7.14	5.73	3.03	3.71	5.35			
	NT2RP2005476	5.22	3.10	3.30	10.18	12.60	10.12	5.36	4.72	5.84	**	+	
	NT2RP2005490	5.25	3.96	4.56	6.13	9.22	5.46	5.31	3.92	5.71			
55	NT2RP2005491	15.97	8.85	12.00	4.52	5.86	4.78	8.53	10.16	9.41	*	-	
	NT2RP2005495	2.68	2.26	2.48	2.05	3.65	3.42	3.01	4.37	2.75			
	NT2RP2005496	9.04	5.08	6.06	16.30	11.28	12.12	9.01	10.34	6.32	*	+	

Table 261

	NT2RP2004709	5.18	3.25	1.93	12.66	12.56	10.94	5.12	4.16	3.85	**	+		
	NT2RP2004710	5.83	4.70	2.80	7.69	7.61	6.76	4.34	3.44	4.54	*	+		
5	NT2RP2004721	11.13	7.44	7.40	6.68	9.65	8.99	11.35	9.52	13.55				
	NT2RP2004736	6.31	5.30	5.26	8.14	9.36	7.77	6.53	5.39	5.85	**	+		
	NT2RP2004743	2.77	1.82	1.65	5.65	6.03	4.15	4.87	6.71	5.76	**	+	**	+
	NT2RP2004750	8.14	5.64	6.27	13.53	14.23	13.30	8.05	8.74	9.81	**	+		
	NT2RP2004755	11.30	7.99	8.26	16.42	20.16	17.92	10.59	13.63	13.47	**	+		
10	NT2RP2004767	6.21	2.89	4.95	9.44	8.05	8.14	4.7	6.19	4.36	*	+		
	NT2RP2004768	9.61	3.95	2.60	2.99	2.03	1.57	2.24	1.57	1.49				
	NT2RP2004775	2.25	2.07	1.48	4.36	5.01	5.07	4.16	3.75	3.44	**	+	**	+
	NT2RP2004791	14.05	7.61	6.73	8.91	10.03	9.17	7.11	6.72	8.15				
	NT2RP2004794	41.53	28.26	27.09	43.02	36.69	32.68	39.95	33.86	41.52				
15	NT2RP2004795	3.77	2.11	2.19	3.89	7.37	3.74	3.78	5.26	5.25		*	+	
	NT2RP2004799	5.43	1.93	3.24	6.30	6.15	4.50	3.93	5.78	3.84				
	NT2RP2004802	4.83	2.53	3.34	7.41	6.03	5.58	2.16	3.27	3.61	*	+		
	NT2RP2004810	3.12	1.86	2.24	8.72	9.56	6.30	5.77	5.46	6.09	**	+	**	+
	NT2RP2004816	4.85	3.14	2.65	6.62	9.96	5.26	6.09	3.65	4.78				
20	NT2RP2004837	13.44	8.28	7.12	11.51	16.25	16.53	19.77	16.72	17.56		*	+	
	NT2RP2004841	2.64	1.81	1.21	3.03	4.37	3.11	1.94	3.01	1.95				
	NT2RP2004847	16.48	11.83	12.45	15.24	18.08	16.57	16.4	14.80	14				
	NT2RP2004861	1.52	1.27	1.44	3.27	3.09	3.21	1.26	1.81	1.52	**	+		
	NT2RP2004897	1.25	0.88	1.99	3.40	2.11	1.91	1.21	2.22	1.75				
25	NT2RP2004932	10.00	7.17	11.03	13.12	14.42	13.51	9.72	9.64	9.65	*	+		
	NT2RP2004933	1.78	1.31	1.88	3.51	3.60	2.84	3.51	3.18	3.33	**	+	**	+
	NT2RP2004936	4.87	2.22	1.77	6.48	8.16	3.31	4.73	2.49	2.48				
	NT2RP2004951	5.43	2.53	1.87	3.02	4.24	3.02	2.87	3.70	11.67				
	NT2RP2004959	8.45	5.08	5.37	8.17	7.86	9.93	4.85	5.55	4.46				
30	NT2RP2004961	5.21	3.54	2.31	7.99	9.20	8.11	4.59	5.46	6.53	**	+		
	NT2RP2004962	4.01	2.64	2.72	5.11	4.60	4.41	3.88	3.76	3.58	*	+		
	NT2RP2004966	2.57	2.53	3.68	2.80	3.88	2.77	2.12	3.33	4.07				
	NT2RP2004967	2.23	2.61	2.86	7.50	6.79	8.12	3.33	4.64	3.83	**	+	*	+
	NT2RP2004974	1.95	1.80	1.93	2.56	3.12	2.39	3.76	2.94	0.71	*	+		
35	NT2RP2004978	6.88	2.95	2.57	5.63	7.09	3.07	4.98	3.62	3.21				
	NT2RP2004982	1.90	1.58	1.47	6.52	6.96	3.08	1.22	2.05	1.93	*	+		
	NT2RP2004985	24.53	11.76	13.37	30.81	35.00	31.74	21.76	19.69	22.43	*	+		
	NT2RP2004999	4.87	3.06	2.28	6.14	7.08	4.89	3.19	4.04	3.16				
	NT2RP2005000	3.68	2.30	2.22	2.75	3.93	3.69	1.87	3.37	3.49				
40	NT2RP2005001	3.57	1.78	2.11	2.93	4.06	3.83	3.59	4.32	2.86				
	NT2RP2005003	4.67	3.07	2.71	7.63	8.71	7.19	5.69	5.10	6.02	**	+	*	+
	NT2RP2005012	6.73	4.06	6.10	5.56	7.80	5.46	4.68	6.28	4.92				
	NT2RP2005018	7.22	3.93	3.53	6.32	10.68	5.74	4.63	4.62	5.01				
	NT2RP2005020	17.60	10.40	7.54	8.46	8.34	5.63	6.22	5.13	5.78				
	NT2RP2005022	4.95	2.69	3.66	5.26	6.40	4.90	4.15	3.47	4.07				
45	NT2RP2005027	22.97	13.64	17.61	9.34	8.66	7.21	22.54	22.24	24.2	*	-		
	NT2RP2005031	1.59	1.04	2.13	1.35	2.05	1.82	1.32	2.27	2.73				
	NT2RP2005035	12.28	9.78	9.98	17.38	24.50	17.61	24.38	23.70	30.68	*	+	**	+
	NT2RP2005037	3.95	3.48	2.80	4.77	7.93	4.42	2.77	3.83	4.79				
	NT2RP2005038	1.07	1.01	1.27	2.71	2.99	1.89	1.22	3.15	1.8	*	+		
50	NT2RP2005048	8.09	4.51	4.12	7.64	8.60	7.45	7.59	5.31	4.79				
	NT2RP2005049	25.41	8.17	11.97	37.61	33.07	31.21	30.69	32.10	38.73	*	+	*	+
	NT2RP2005073	4.93	2.00	2.06	7.13	4.92	3.75	2.76	2.91	4.07				
	NT2RP2005077	4.59	2.92	2.93	3.87	3.63	3.16	2.4	2.61	2.69				
	NT2RP2005108	3.21	2.75	1.61	3.23	2.96	2.92	1.57	2.62	2.37				
	NT2RP2005116	9.11	5.71	5.87	6.08	9.75	7.92	7.26	7.17	8.23				
55	NT2RP2005126	8.28	8.63	9.53	6.69	10.50	10.65	4.18	6.96	4.15		*	-	
	NT2RP2005135	3.79	3.03	2.85	3.91	5.50	2.16	3.03	4.16	3.27				

Table 260

	NT2RP2004242	4.01	3.66	4.18	4.80	6.97	3.56	2.91	4.22	3.72			
	NT2RP2004245	4.75	2.29	3.26	4.55	5.39	2.63	3.01	2.48	2.79			
5	NT2RP2004270	18.23	8.30	7.67	19.68	17.41	13.31	11.69	12.72	8.05			
	NT2RP2004300	3.69	2.58	2.90	3.43	6.04	3.65	2.47	3.40	4.86			
	NT2RP2004384	6.67	2.88	6.27	10.77	12.81	11.19	6.73	6.65	7.62	**	+	
	NT2RP2004313	3.69	3.44	2.33	4.32	4.99	4.51	2.56	4.15	4.27	*	+	
	NT2RP2004316	4.16	1.43	2.32	4.51	4.31	4.04	2.43	3.50	4.17			
10	NT2RP2004321	15.92	11.27	11.28	36.60	56.46	33.80	10.57	12.49	9.91	*	+	
	NT2RP2004336	2.22	1.97	1.95	1.98	2.72	1.41	1.95	2.65	2.2			
	NT2RP2004339	18.02	10.18	7.42	25.42	25.92	20.21	14.54	10.64	9.51	*	+	
	NT2RP2004347	6.36	3.28	2.51	3.98	5.62	3.33	2.91	2.22	3.77			
	NT2RP2004364	7.25	3.84	3.16	7.45	10.83	6.50	5.33	5.38	5.14			
15	NT2RP2004365	3.92	1.67	1.92	3.47	3.94	3.44	1.64	2.60	3.66			
	NT2RP2004366	3.77	1.94	2.27	3.01	4.43	2.63	2.6	3.92	2.96			
	NT2RP2004373	2.38	2.55	1.79	5.73	5.73	2.95	2.28	3.83	3.83			
	NT2RP2004375	14.49	9.73	10.51	9.34	13.60	9.23	5.43	7.02	8.38			
	NT2RP2004389	6.54	5.30	4.58	4.64	5.83	5.40	4.4	4.73	4.62			
20	NT2RP2004392	28.46	15.89	13.93	32.21	29.99	20.99	14.28	13.07	11.38			
	NT2RP2004396	12.58	7.77	8.62	10.01	8.33	7.76	2.74	2.93	6		*	-
	NT2RP2004399	7.37	3.73	4.44	6.18	6.63	5.28	3.66	5.23	7.06			
	NT2RP2004400	3.45	1.87	1.89	5.43	5.79	4.47	2.84	3.98	3.76	*	+	
	NT2RP2004404	11.50	7.62	6.89	11.66	13.80	10.35	8.27	8.35	9.19			
25	NT2RP2004410	11.23	11.38	11.20	17.64	15.77	17.12	11.2	18.45	13.95	**	+	
	NT2RP2004412	4.89	2.82	3.13	4.05	4.86	3.06	2.32	3.89	3.43			
	NT2RP2004414	6.08	2.18	5.00	3.14	3.56	2.80	1.59	3.95	2.41			
	NT2RP2004425	2.01	1.60	1.70	2.43	4.37	2.34	2.53	1.37	3.45			
	NT2RP2004447	3.57	2.63	1.82	4.60	4.54	3.34	3.94	3.07	2.46			
30	NT2RP2004463	11.21	7.40	6.24	12.62	8.89	9.28	9.29	8.97	10.07			
	NT2RP2004476	4.90	3.15	2.20	5.47	5.87	6.15	2.61	3.85	5.36	*	+	
	NT2RP2004488	5.90	4.58	3.55	4.28	5.12	3.55	2.91	3.17	2.6			
	NT2RP2004490	4.32	3.15	2.55	3.51	4.12	4.44	2.62	3.95	8.62			
	NT2RP2004495	12.24	5.83	8.88	11.24	10.73	8.49	9.47	11.08	18.95			
35	NT2RP2004512	5.33	2.48	2.45	3.28	4.26	3.70	3.48	2.44	3.06			
	NT2RP2004523	10.16	5.01	3.79	10.11	8.70	10.80	6.51	6.83	6.35			
	NT2RP2004524	3.86	3.51	2.47	5.08	4.81	4.16	5.08	3.55	3.98	*	+	
	NT2RP2004536	11.38	9.71	7.82	9.14	12.16	9.03	6.49	7.82	9.84			
	NT2RP2004538	38.06	30.58	30.32	62.14	68.91	71.97	40.07	32.51	41.6	**	+	
	NT2RP2004548	5.50	4.46	3.74	10.83	12.12	12.54	4.81	5.53	8.83	**	+	
40	NT2RP2004551	3.34	1.83	3.26	4.96	5.20	3.92	2.98	2.39	13.41	*	+	
	NT2RP2004556	8.58	7.04	6.71	15.74	11.77	13.75	8.42	10.11	9.77	**	+	
	NT2RP2004568	19.23	11.22	8.88	15.82	10.64	12.31	13.87	9.13	9.97			
	NT2RP2004580	7.17	4.71	2.64	11.67	9.76	7.99	7.1	5.36	6.59	*	+	
45	NT2RP2004585	10.92	6.41	6.18	10.89	10.92	9.49	8.51	7.18	15.75			
	NT2RP2004587	2.30	1.65	0.84	2.47	1.80	1.78	1.76	1.48	2.46			
	NT2RP2004594	5.87	5.87	4.84	5.34	8.13	4.27	3.88	5.17	7.53			
	NT2RP2004600	1.88	2.05	1.13	2.29	2.11	2.15	1.86	2.50	1.01			
	NT2RP2004602	4.95	4.31	4.04	9.75	8.80	8.23	5.05	5.03	6.56	**	+	
	NT2RP2004606	11.77	11.03	6.62	13.49	15.68	9.80	18.35	17.80	17.2		**	+
50	NT2RP2004614	7.71	4.83	3.32	3.55	3.54	4.41	5.21	3.79	4.38			
	NT2RP2004648	6.00	3.54	2.10	4.35	4.65	3.29	4.52	3.41	7.33			
	NT2RP2004655	13.74	9.02	8.53	4.81	7.79	4.98	3.5	4.03	6.04		*	-
	NT2RP2004664	6.11	4.51	4.83	11.64	9.59	6.61	6.34	6.03	5.86			
	NT2RP2004670	3.00	2.33	2.81	3.65	3.96	4.05	3.26	3.38	4.44	**	+	
55	NT2RP2004675	5.69	3.74	5.15	11.77	11.28	10.21	5.45	4.84	5.26	**	+	
	NT2RP2004681	5.04	3.65	4.78	6.53	9.83	7.35	6.37	6.72	6.63	*	+	**
	NT2RP2004689	2.24	1.16	1.68	3.06	4.60	6.68	1.3	2.45	1.72	*	+	

Table 259

	NT2RP2003825	9.16	5.63	6.57	17.27	18.54	12.04	6.67	8.08	14.03	*	+		
	NT2RP2003840	10.64	4.89	5.66	8.31	7.78	5.93	7.12	5.91	8.06				
5	NT2RP2003857	12.72	6.86	6.25	8.31	8.84	9.18	7.95	6.05	8.74				
	NT2RP2003859	6.93	3.73	2.73	12.12	10.40	13.45	5.71	3.90	6.36	**	+		
	NT2RP2003871	3.42	3.01	2.13	9.67	10.18	8.65	5.24	4.53	5.97	**	+	*	+
	NT2RP2003876	7.74	4.51	4.43	5.67	8.07	7.43	4.37	5.53	5.6				
	NT2RP2003878	4.47	2.22	2.10	3.89	4.71	3.64	3.95	3.56	4.06				
10	NT2RP2003885	5.69	2.59	2.76	3.73	7.92	5.39	4.25	4.87	6.01				
	NT2RP2003898	10.09	7.67	7.33	11.75	12.18	9.75	5.01	8.03	5.65				
	NT2RP2003902	10.41	8.37	6.78	8.14	9.71	9.88	7.68	5.42	8.06				
	NT2RP2003912	13.81	9.98	7.42	16.63	17.90	13.52	10.9	14.66	13.18				
	NT2RP2003931	3.74	1.68	1.44	2.28	2.88	2.54	2.24	1.94	2.65				
15	NT2RP2003940	18.24	10.75	11.51	44.72	39.79	24.81	16	14.58	19.02	*	+		
	NT2RP2003950	3.98	2.45	3.31	3.52	4.06	3.60	3	3.05	3.52				
	NT2RP2003952	5.00	3.18	4.24	4.00	4.74	3.20	2.55	2.62	4.33				
	NT2RP2003968	13.52	6.81	6.24	9.83	14.58	9.98	4.25	4.49	10.21				
	NT2RP2003976	5.76	3.40	2.77	10.86	15.30	22.19	5.6	7.99	7.6	*	+		
20	NT2RP2003981	5.81	3.89	2.20	4.65	4.94	4.43	4.88	3.67	4.28				
	NT2RP2003984	11.22	7.15	6.30	8.47	13.43	9.96	9.18	9.47	16.24				
	NT2RP2003986	11.50	5.47	4.61	14.29	15.56	15.24	7.95	7.99	8.32	*	+		
	NT2RP2003988	5.84	4.44	3.08	11.21	13.07	8.96	7.35	4.72	6.91	**	+		
	NT2RP2004013	19.46	11.40	12.00	20.33	26.92	19.32	8.59	11.59	12.13				
25	NT2RP2004014	5.88	5.77	8.06	11.00	14.73	13.84	6.02	5.49	4.74	**	+		
	NT2RP2004036	4.76	2.41	3.64	4.63	4.19	5.70	3.7	3.95	3.26				
	NT2RP2004041	2.79	3.61	3.30	4.01	6.06	4.15	3.2	4.29	4.43				
	NT2RP2004042	4.23	3.45	2.82	4.59	3.59	5.00	3.97	2.94	3.64				
	NT2RP2004049	5.52	3.09	3.20	5.68	4.82	4.18	3.14	3.78	3.4				
30	NT2RP2004060	6.54	4.19	4.75	5.31	7.44	5.90	6.84	5.31	6.57				
	NT2RP2004066	7.62	3.57	3.11	8.07	8.17	6.09	3.54	4.23	4.08				
	NT2RP2004069	2.46	2.35	2.84	3.73	4.30	3.52	3.02	4.14	4.07	**	+	*	+
	NT2RP2004076	1.40	1.15	1.26	2.49	2.65	1.93	1.27	2.46	1.33	**	+		
	NT2RP2004080	2.70	2.23	2.55	3.88	5.93	4.96	4.18	5.58	4.25	*	+	*	+
	NT2RP2004081	2.74	2.99	2.36	3.72	4.51	3.72	1.45	3.28	1.61	*	+		
35	NT2RP2004098	10.83	5.42	4.87	10.62	9.37	7.52	6.04	4.69	6.05				
	NT2RP2004108	15.24	8.74	6.82	24.00	21.97	22.21	10.22	12.30	14.43	**	+		
	NT2RP2004124	5.29	4.13	3.63	5.87	5.42	5.25	4.18	2.84	4.23				
	NT2RP2004130	9.77	7.17	7.05	9.85	13.14	10.78	12.57	13.32	11.04		*	+	
40	NT2RP2004133	11.24	7.82	7.31	10.46	12.30	8.54	8.71	9.42	8.83				
	NT2RP2004141	4.33	2.78	3.55	5.05	6.27	4.10	3.83	4.25	5.14				
	NT2RP2004142	3.53	1.25	3.26	3.70	5.10	5.11	2.84	4.94	3.66				
	NT2RP2004152	2.68	1.78	2.43	4.24	5.04	5.23	2.05	2.34	1.5	**	+		
	NT2RP2004165	21.03	8.19	8.39	7.87	8.05	7.98	5.38	6.52	6.22				
	NT2RP2004170	7.13	4.37	2.78	6.23	7.89	6.07	5.24	5.06	3.73				
45	NT2RP2004172	3.69	2.25	1.50	2.50	3.71	2.71	2.83	3.52	1.97				
	NT2RP2004176	7.84	4.13	3.67	5.48	5.12	4.33	5.56	7.38	6.12				
	NT2RP2004179	6.87	2.52	2.41	5.35	4.30	3.84	3.98	4.72	4.2				
	NT2RP2004187	3.69	2.64	1.86	5.62	6.94	5.86	3.38	4.90	4.03	**	+		
	NT2RP2004190	2.07	2.03	2.45	3.29	3.28	2.78	5.06	5.55	4.18	*	+	**	+
50	NT2RP2004194	6.67	3.78	5.18	7.29	8.60	7.46	5.61	6.42	7				
	NT2RP2004196	20.28	5.85	8.55	16.34	14.05	15.75	7.78	7.99	8.4				
	NT2RP2004205	10.63	6.42	6.10	11.21	13.23	11.22	6.53	6.15	7.63				
	NT2RP2004207	4.42	3.24	2.70	3.44	4.24	3.84	3.13	3.26	3.82				
	NT2RP2004226	4.97	4.89	4.35	4.76	5.20	4.65	3.73	3.67	3.35		**	-	
	NT2RP2004232	2.49	1.77	2.98	3.76	4.69	3.30	2.85	3.10	2.15	*	+		
55	NT2RP2004239	4.49	3.56	3.79	6.17	7.37	6.14	4.15	5.46	4.58	**	+		
	NT2RP2004240	6.30	3.45	4.77	13.34	11.74	9.18	6.02	6.36	6.66	*	+		

Table 258

	NT2RP2003470	11.59	7.42	9.22	28.44	23.50	24.05	11.29	12.07	8.19	**	+		
	NT2RP2003471	0.69	0.28	0.53	1.86	1.08	1.71	2.23	2.31	0.86	*	+		
5	NT2RP2003480	15.63	7.31	7.47	13.91	14.92	13.14	9.58	7.59	11				
	NT2RP2003495	6.78	5.33	-4.65	5.96	5.20	6.08	4.27	5.58	4.14				
	NT2RP2003499	3.16	1.30	1.31	2.42	1.62	2.16	3.79	4.26	2.53				
	NT2RP2003505	2.95	2.52	1.64	4.06	3.25	3.65	2.65	3.70	2.81	*	+		
	NT2RP2003506	4.36	2.44	2.89	4.61	6.57	3.32	3.86	4.37	5.74				
10	NT2RP2003511	5.80	4.98	5.36	9.63	8.04	5.73	6.43	6.77	8.36		*	+	
	NT2RP2003513	3.23	2.52	3.10	3.94	3.00	3.76	2.27	3.48	3.18				
	NT2RP2003517	1.52	0.95	2.01	2.87	2.13	1.37	2.66	3.16	3.17		*	+	
	NT2RP2003522	21.16	8.31	12.55	21.51	17.78	15.40	9.2	5.69	8.01				
	NT2RP2003525	6.58	6.05	5.00	12.44	12.64	12.83	8.86	7.54	7.95	**	+	*	+
15	NT2RP2003533	7.73	4.59	4.51	11.94	12.52	10.34	6.62	8.25	8.72	**	+		
	NT2RP2003541	9.89	7.73	6.72	8.34	7.49	6.40	6.78	6.83	5.85				
	NT2RP2003543	4.46	3.26	2.49	5.01	7.76	4.19	6.57	7.85	7.39		**	+	
	NT2RP2003545	6.37	3.24	4.48	2.58	2.60	1.05	1.96	3.63	2.3				
	NT2RP2003559	1.78	1.16	2.25	3.59	3.08	3.14	2.24	2.88	3.16	*	+		
20	NT2RP2003564	1.65	1.70	1.81	2.44	3.74	2.88	2.97	3.23	1.66	*	+		
	NT2RP2003565	9.14	3.08	4.12	8.63	10.17	6.24	4.03	4.24	3.56				
	NT2RP2003567	7.44	5.21	4.96	7.20	9.00	7.04	7.75	6.53	4.86				
	NT2RP2003575	5.24	1.86	2.00	2.78	2.67	1.70	1.73	2.24	4.67				
	NT2RP2003576	208.36	132.21	112.56	100.63	118.10	86.36	71.48	50.82	50.69		*	-	
25	NT2RP2003579	56.28	38.17	48.67	28.49	15.58	24.16	19.34	17.93	21.34	*	-	**	-
	NT2RP2003581	4.71	3.22	3.45	3.09	5.04	4.47	3.46	3.82	4.77				
	NT2RP2003587	8.55	4.99	7.99	8.79	9.50	8.44	7.38	8.78	13.4				
	NT2RP2003590	11.27	7.70	8.07	4.15	4.86	4.77	3.73	6.36	4.84	*	-	*	-
	NT2RP2003593	9.63	4.82	5.47	13.80	9.75	5.79	6.89	8.08	6.91				
	NT2RP2003596	3.20	2.89	2.89	6.00	8.78	7.99	4.62	4.90	7.08	**	+	*	+
30	NT2RP2003599	8.81	5.81	5.81	8.37	10.49	10.48	10.61	8.00	12.61				
	NT2RP2003600	3.15	1.54	2.36	3.63	5.05	4.21	2.91	3.54	3.28	*	+		
	NT2RP2003604	8.61	4.63	5.27	5.66	7.11	7.00	5.84	5.70	5.33				
	NT2RP2003629	0.93	0.41	0.97	1.80	1.56	1.57	0.76	2.29	1.4	*	+		
35	NT2RP2003630	3.31	2.56	2.95	6.23	8.50	6.34	5.52	5.72	4.54	**	+	**	+
	NT2RP2003643	16.50	10.48	12.66	12.59	15.91	12.75	9.42	11.38	10.03				
	NT2RP2003655	4.54	2.17	1.95	4.91	4.47	3.19	4.31	4.99	6.38				
	NT2RP2003664	7.29	4.58	3.44	9.78	13.11	10.33	7.53	12.65	18.19	*	+		
	NT2RP2003668	7.64	3.93	2.99	7.77	11.11	7.27	3.61	4.49	4.92				
	NT2RP2003687	3.50	2.00	2.53	2.44	3.28	2.52	1.34	3.20	1.86				
40	NT2RP2003691	3.51	2.23	2.36	4.83	5.26	4.14	2.6	3.93	3.34	*	+		
	NT2RP2003702	4.72	3.23	2.91	5.75	5.42	5.03	3.29	5.65	2.48	*	+		
	NT2RP2003704	3.03	1.02	1.33	3.00	4.19	2.96	1.48	4.19	2.8				
	NT2RP2003706	0.54	0.54	0.40	1.92	1.23	0.53	1.37	2.50	2.1		*	+	
	NT2RP2003713	3.77	2.04	1.68	4.89	3.40	3.69	3.54	1.79	2.29				
45	NT2RP2003714	16.93	11.05	8.85	15.34	13.25	10.73	6.94	5.43	4.92				
	NT2RP2003727	9.17	5.59	4.98	8.92	8.98	7.11	5.82	4.15	6.45				
	NT2RP2003737	4.49	2.62	2.06	3.80	4.50	3.26	2.92	3.29	5.35				
	NT2RP2003751	0.82	0.97	1.07	1.33	1.62	0.98	1.33	0.88	0.72				
	NT2RP2003760	3.61	2.60	1.42	4.28	5.22	4.19	4.75	3.97	7.45	*	+		
50	NT2RP2003764	4.43	3.65	3.32	3.81	3.64	3.20	3.86	3.12	8.26				
	NT2RP2003769	3.03	1.62	1.45	3.28	5.14	3.51	3.96	2.62	2.26				
	NT2RP2003770	11.88	6.14	5.72	10.96	9.10	9.34	9.86	5.90	7.19				
	NT2RP2003777	8.28	5.95	4.45	12.14	8.07	7.31	6.16	4.05	5.91				
	NT2RP2003781	6.93	4.17	4.88	6.60	9.83	10.25	6.27	5.64	6.39				
55	NT2RP2003785	5.07	3.24	3.30	5.65	5.69	5.57	6.33	7.73	14.42	*	+		
	NT2RP2003793	9.26	6.02	4.92	6.26	7.16	5.41	4.28	4.76	5.16				
	NT2RP2003806	6.44	4.78	6.02	12.68	12.04	12.13	5.52	7.88	5.99	**	+		

Table 257

	NT2RP2003129	3.68	2.64	1.93	5.72	5.89	5.75	3.03	4.40	2.82	**	+		
5	NT2RP2003137	2.40	2.79	2.71	6.74	6.38	5.76	4.22	6.41	4.31	**	+	*	+
	NT2RP2003138	6.42	2.67	2.97	5.99	6.92	3.98	5.12	3.06	1.92				
	NT2RP2003146	4.44	2.51	1.78	3.73	3.26	2.77	3.76	2.57	1.66				
	NT2RP2003148	9.10	6.45	5.51	11.73	13.86	11.19	8.71	8.13	7.46	*	+		
	NT2RP2003150	3.26	2.20	1.35	8.65	2.99	4.86	3.92	2.84	8.35				
10	NT2RP2003157	7.49	3.86	3.67	8.41	10.43	9.55	4.96	6.45	5.87	*	+		
	NT2RP2003158	1.98	1.89	2.17	2.26	3.00	2.46	2.43	2.76	2.85			*	+
	NT2RP2003161	1.04	1.33	0.76	2.12	4.38	4.18	1.59	2.84	8.91	*	+		
	NT2RP2003164	2.83	1.78	1.70	2.90	2.78	2.57	2.53	2.97	2.44				
	NT2RP2003165	4.31	2.10	2.06	5.98	4.84	6.84	5.12	3.81	4.72	*	+		
	NT2RP2003177	3.18	2.52	2.22	3.53	2.99	3.63	4.35	2.80	2.79				
15	NT2RP2003179	4.54	3.39	3.36	5.90	7.70	7.29	4.85	4.79	6.24	**	+		
	NT2RP2003194	16.94	9.59	9.74	7.86	8.77	6.84	7.23	6.50	9.93				
	NT2RP2003206	0.19	0.73	0.54	2.02	2.10	1.11	1.07	1.15	1.17	*	+	*	+
	NT2RP2003210	5.52	2.50	2.65	2.94	4.61	3.60	3.44	3.99	4.15				
	NT2RP2003227	2.55	1.52	2.78	3.96	4.66	3.48	2.52	3.60	4.44	*	+		
20	NT2RP2003228	5.50	4.11	4.96	4.07	4.64	3.51	3.63	3.86	2.66				
	NT2RP2003230	1.04	1.41	1.38	3.75	3.72	3.44	8.77	4.96	7.21	**	+	**	+
	NT2RP2003231	6.83	5.52	4.87	9.61	7.64	6.47	5.75	5.89	8.09				
	NT2RP2003237	4.46	2.56	2.35	5.51	7.13	6.33	3.56	4.31	3.67	*	+		
	NT2RP2003239	4.50	2.01	3.71	6.44	6.32	5.76	4.01	4.23	4.42	*	+		
25	NT2RP2003243	5.46	3.20	3.57	7.44	6.11	7.58	5.91	6.40	3.87	*	+		
	NT2RP2003245	5.61	3.24	3.60	7.47	8.92	7.01	5.38	4.10	6.74	*	+		
	NT2RP2003267	3.97	3.06	3.71	7.15	8.86	6.88	4.28	4.40	5.84	**	+		
	NT2RP2003272	5.37	3.98	5.63	6.49	6.56	6.62	7.54	6.51	7.61	*	+	*	+
	NT2RP2003277	9.14	5.91	4.66	7.52	10.35	9.11	9.97	7.77	15.8				
30	NT2RP2003280	3.01	2.25	1.41	4.02	6.71	7.68	6.13	4.20	7.59	*	+	*	+
	NT2RP2003286	3.53	1.84	2.37	2.62	3.15	2.83	2.96	2.70	4.01				
	NT2RP2003293	6.85	4.64	6.03	12.22	12.54	11.97	6.66	5.15	8.8	**	+		
	NT2RP2003295	4.81	3.25	3.18	3.96	8.36	5.27	4.16	4.98	3				
	NT2RP2003297	1.97	1.06	1.42	2.82	3.09	2.49	1.97	1.89	1.68	*	+		
35	NT2RP2003300	5.99	4.89	4.68	7.75	7.40	7.47	7.28	9.19	9.08	**	+	*	+
	NT2RP2003302	4.65	3.24	4.39	8.90	10.20	7.29	4.36	7.27	5.11	**	+		
	NT2RP2003307	1.67	1.09	0.57	2.24	1.67	2.40	2.82	1.84	1.76				
	NT2RP2003308	3.09	2.17	1.85	4.09	5.19	2.83	3.04	2.74	3.16				
	NT2RP2003311	6.85	3.58	2.13	4.65	6.66	4.36	3.88	3.65	4.23				
	NT2RP2003329	3.07	1.86	1.87	3.19	5.07	3.49	3.77	3.82	5.96				
40	NT2RP2003339	2.38	1.55	1.29	2.90	3.98	3.91	2.69	3.47	2.24	*	+		
	NT2RP2003345	1.83	1.44	1.40	1.51	1.52	1.92	2.28	2.65	1.28				
	NT2RP2003347	1.48	2.10	1.67	2.03	5.75	1.76	2.44	3.10	4.09			*	+
	NT2RP2003367	1.26	0.98	1.42	1.39	1.59	1.55	1.21	2.14	1.04				
	NT2RP2003369	3.82	2.31	1.37	1.62	2.10	1.87	3.19	2.85	1.99				
45	NT2RP2003383	7.18	3.57	4.41	16.30	14.96	15.98	8.79	9.62	11.29	**	+	*	+
	NT2RP2003390	9.92	6.14	6.73	11.71	12.19	9.52	7.92	9.43	8.34				
	NT2RP2003391	35.23	21.64	23.50	36.95	36.23	27.51	23.69	17.29	17.85				
	NT2RP2003393	2.40	1.57	1.83	4.13	5.18	3.56	3.96	4.34	3.87	*	+	**	+
	NT2RP2003394	4.02	2.41	2.76	12.16	9.99	10.68	6.12	6.15	3.96	**	+		
50	NT2RP2003401	2.33	1.80	1.86	3.02	4.68	2.41	3.02	4.51	3.57			*	+
	NT2RP2003403	1.23	1.40	1.41	3.20	3.23	4.51	3.04	3.80	3.41	**	+	**	+
	NT2RP2003433	8.96	4.52	3.52	6.71	5.66	5.39	7.4	6.01	5.01				
	NT2RP2003445	3.20	3.09	2.41	6.94	6.16	6.94	13.01	11.43	14.04	**	+	**	+
	NT2RP2003446	5.05	4.02	2.72	4.09	6.31	3.82	5.45	4.95	5.35				
	NT2RP2003456	4.21	2.96	2.69	10.80	8.14	8.43	6.15	5.44	4.71	**	+	*	+
55	NT2RP2003466	5.26	3.68	3.82	5.95	5.44	4.60	3.82	5.23	9				
	NT2RP2003469	3.53	2.12	2.45	3.89	4.69	5.28	2.75	4.01	3.09	*	+		

Table 256

5	NT2RP2002734	4.55	3.02	5.80	12.41	12.84	10.69	6.86	8.07	7.85	**	+	"	+
	NT2RP2002736	3.63	2.27	2.67	2.07	2.02	2.04	2.87	2.60	2.01				
	NT2RP2002740	2.59	1.02	0.94	3.18	2.63	2.29	2.78	2.96	1.96				
	NT2RP2002741	5.52	4.27	3.15	7.73	8.99	8.94	4.51	5.06	7.43	**	+		
	NT2RP2002750	7.28	6.29	4.77	14.35	17.57	18.80	8.32	9.26	7.61	**	+		
	NT2RP2002752	11.68	7.46	7.74	12.78	17.74	15.50	11.31	10.02	12.22	*	+		
10	NT2RP2002753	11.55	5.48	11.53	10.53	6.13	11.57	7.42	7.93	9.43				
	NT2RP2002760	8.78	4.40	4.62	7.89	8.63	6.01	6.34	6.38	7.33				
	NT2RP2002769	3.29	2.63	2.68	3.72	6.64	6.67	2.86	4.11	3.55	*	+		
	NT2RP2002778	9.07	6.03	9.70	7.44	6.87	7.92	6.93	7.76	4.98				
	NT2RP2002791	6.58	4.82	4.00	9.50	14.75	9.25	8.23	6.79	7.02	*	+		
15	NT2RP2002800	6.57	4.20	5.63	10.46	11.33	12.38	5.4	8.07	7.04	**	+		
	NT2RP2002805	1.48	1.18	0.66	2.57	1.66	1.18	2.89	3.53	1.96			"	+
	NT2RP2002811	5.70	5.54	4.77	8.54	7.13	7.69	6.53	7.67	6.08	**	+		
	NT2RP2002824	9.12	5.93	7.91	13.68	13.22	9.65	9.82	10.22	11.6	*	+		
	NT2RP2002839	3.89	2.03	2.96	3.87	4.52	3.28	3.17	3.43	3.41				
20	NT2RP2002845	2.29	1.84	1.77	4.04	4.31	4.72	3.6	4.26	3.16	**	+	**	+
	NT2RP2002857	0.99	1.45	1.80	1.98	2.27	1.76	2.36	3.14	1.89				
	NT2RP2002862	11.21	6.20	5.58	10.84	12.86	10.44	6.99	7.12	10.71				
	NT2RP2002880	5.70	4.03	2.74	3.50	4.84	3.87	4.05	5.72	5				
	NT2RP2002885	6.90	4.59	4.82	5.83	6.45	4.16	3.34	4.76	3.08				
25	NT2RP2002891	5.76	3.80	3.33	5.44	6.69	6.13	4.92	4.49	5.35				
	NT2RP2002907	4.12	1.98	2.30	4.77	3.91	2.49	2.25	3.24	2.04				
	NT2RP2002925	3.23	2.04	2.18	4.98	4.44	5.21	3.38	2.81	4.67	**	+		
	NT2RP2002927	14.45	8.55	11.84	14.25	14.86	13.10	10.66	9.50	13.04				
	NT2RP2002928	1.42	1.26	2.32	3.26	2.52	3.14	1.44	1.91	1.88	*	+		
30	NT2RP2002929	6.54	3.13	3.18	6.60	7.00	5.63	5.25	5.85	5.87				
	NT2RP2002934	5.87	2.70	3.00	3.46	2.95	4.09	3.58	3.88	3.47				
	NT2RP2002939	6.87	3.02	3.14	4.78	4.45	4.28	3.95	4.36	3.63				
	NT2RP2002942	4.16	2.79	3.25	6.95	8.21	6.01	4.14	5.76	4.58	**	+		
	NT2RP2002954	3.73	2.07	3.02	3.75	4.03	3.04	2.28	3.89	5.22				
	NT2RP2002959	5.43	4.36	4.62	6.19	7.91	6.08	3.63	5.75	5.03	*	+		
35	NT2RP2002974	2.77	2.53	1.82	5.32	4.88	3.20	3.66	3.70	3.24	*	+	"	+
	NT2RP2002976	1.81	1.66	2.46	4.07	3.02	2.77	2.16	2.65	2.13	*	+		
	NT2RP2002979	10.96	6.09	6.26	13.05	14.90	10.76	8.18	9.68	7.32				
	NT2RP2002980	8.71	5.49	6.33	14.65	15.05	11.66	8.24	9.16	9.26	**	+		
	NT2RP2002986	8.28	6.07	5.22	8.21	6.48	6.46	9.09	7.74	9.39				
40	NT2RP2002987	6.13	3.28	3.28	8.77	8.51	7.89	4.85	7.00	9.15	*	+		
	NT2RP2002988	34.52	23.01	24.20	21.24	19.88	21.98	15.82	15.65	16.56			"	-
	NT2RP2002993	4.35	3.19	4.08	2.57	3.44	2.83	3.21	3.84	2.8				
	NT2RP2003000	6.81	5.24	5.01	12.83	14.50	14.13	6.77	6.65	8.42	**	+		
	NT2RP2003006	3.03	1.86	2.21	2.77	3.21	3.26	2.46	3.49	5.58				
45	NT2RP2003020	7.91	3.15	3.03	14.51	13.63	11.55	10.67	9.71	9.8	**	+	"	+
	NT2RP2003032	4.25	3.36	3.04	5.65	7.30	4.26	5.14	2.86	5.02				
	NT2RP2003034	8.64	4.19	5.82	12.73	13.68	11.86	9.6	7.30	8.21	**	+		
	NT2RP2003042	3.77	2.17	2.53	3.68	4.54	3.65	3.09	3.66	3.89				
	NT2RP2003050	2.09	1.93	2.12	2.58	4.04	3.16	2.04	3.12	2.84	*	+		
50	NT2RP2003060	6.89	6.04	6.20	6.11	6.61	6.02	4.64	5.08	4.58			**	-
	NT2RP2003073	5.10	4.79	4.81	10.73	11.79	9.58	6.83	8.25	4.87	**	+		
	NT2RP2003099	3.77	3.26	2.86	5.64	6.07	7.01	4.27	4.57	5.19	**	+	"	+
	NT2RP2003106	3.73	1.70	0.71	4.43	4.78	3.41	2.53	3.13	3.98				
	NT2RP2003115	12.63	7.03	6.49	10.94	5.30	4.75	6.51	5.99	10.91				
55	NT2RP2003117	9.96	4.65	5.66	15.04	15.48	11.38	8.83	7.84	6.17	*	+		
	NT2RP2003121	3.53	2.40	1.92	4.30	5.00	3.18	3.72	4.52	4.42			"	+
	NT2RP2003125	5.32	2.20	2.34	3.41	4.18	3.51	3.6	4.31	3.11				
	NT2RP2003127	3.09	3.27	3.35	3.25	3.63	2.74	2.3	4.46	3.68				

Table 255

	NT2RP2002292	13.36	8.93	10.00	7.24	12.33	7.03	8.51	6.90	8.43				
5	NT2RP2002299	4.86	3.21	3.87	7.31	5.99	7.44	5.79	6.94	6.46	*	+	*	+
	NT2RP2002304	3.12	1.09	1.07	3.72	6.64	4.48	2.39	2.10	2.14	*	+		
	NT2RP2002312	3.00	2.02	1.91	4.87	5.25	3.26	3.11	3.70	3.89	*	+	*	+
	NT2RP2002316	2.57	2.29	2.38	6.74	6.43	5.78	3.25	3.23	4.39	**	+	*	+
	NT2RP2002325	2.17	2.03	1.50	3.32	3.39	2.92	1.65	3.11	3.18	**	+		
10	NT2RP2002333	6.45	4.83	4.75	7.88	10.32	7.81	5.66	5.80	6.3	*	+		
	NT2RP2002371	4.90	4.23	3.63	9.29	8.56	8.25	9.75	10.58	7.26	**	+	**	+
	NT2RP2002373	5.37	4.02	2.70	5.83	10.05	6.25	5.7	8.27	6.72				
	NT2RP2002381	0.73	0.29	0.85	0.79	0.90	2.57	1.16	2.65	1.41				
	NT2RP2002385	7.34	2.40	2.24	6.24	3.86	3.39	5.09	3.89	4.74				
	NT2RP2002394	1.71	0.33	0.18	1.03	1.49	1.31	0.28	1.27	2.19				
15	NT2RP2002408	2.38	1.66	1.45	4.45	2.73	2.67	1.95	4.44	3.16				
	NT2RP2002409	29.85	16.62	15.12	29.12	39.51	28.40	19.16	20.28	16.59				
	NT2RP2002424	3.78	2.45	1.98	3.14	4.67	3.25	3.81	5.82	3.46				
	NT2RP2002426	5.16	3.36	3.05	8.68	9.29	8.07	5.5	8.86	7.03	**	+	*	+
	NT2RP2002429	6.36	5.02	5.09	9.72	12.33	8.37	9.84	17.67	16.81	*	+	*	+
20	NT2RP2002437	3.49	2.56	3.29	4.17	7.17	4.10	3.26	6.17	5.32				
	NT2RP2002439	11.07	5.27	5.30	11.81	8.46	7.22	11.52	9.36	7.78				
	NT2RP2002442	6.40	2.74	3.03	4.62	5.05	4.46	4.75	2.98	3.74				
	NT2RP2002457	2.28	2.49	1.70	3.54	4.01	3.48	4.07	3.72	3.08	**	+	*	+
	NT2RP2002464	5.19	2.78	3.13	3.90	4.79	4.00	5.08	3.74	4				
25	NT2RP2002475	3.58	3.74	3.05	8.04	7.22	4.99	7.48	6.02	7.62	*	+	**	+
	NT2RP2002479	3.49	2.33	2.32	3.60	4.32	2.72	2.92	2.66	5.14				
	NT2RP2002487	4.86	2.73	2.49	4.04	4.25	4.00	3.16	3.11	3.07				
	NT2RP2002498	2.48	0.99	1.21	3.47	2.96	2.55	1.35	1.52	1.58				
	NT2RP2002503	13.02	6.05	8.78	12.14	16.89	12.87	9.04	8.81	7.66				
30	NT2RP2002504	6.63	3.00	4.84	4.05	6.27	4.67	6.68	4.71	5.18				
	NT2RP2002510	15.40	9.87	11.00	12.38	17.28	17.15	18.56	12.92	13.19				
	NT2RP2002520	1.61	1.78	1.33	4.08	3.77	4.83	3.97	4.73	4.31	**	+	**	+
	NT2RP2002527	11.26	7.87	9.14	12.36	15.57	11.93	8.08	6.87	9.06				
	NT2RP2002533	15.80	10.32	13.55	16.21	16.47	14.65	18.71	12.94	18.73				
	NT2RP2002537	6.78	4.47	5.46	7.12	8.21	8.66	4.34	3.85	6.54	*	+		
35	NT2RP2002542	11.84	6.86	7.87	24.97	24.70	21.27	12.25	9.81	10.65	**	+		
	NT2RP2002546	3.51	1.75	1.39	2.49	2.71	2.52	4.4	3.54	3.7				
	NT2RP2002549	8.05	4.99	5.19	5.57	6.51	7.45	6.2	3.49	5.35				
	NT2RP2002564	13.08	7.54	8.36	11.61	12.09	10.41	11.1	8.10	13.89				
	NT2RP2002591	9.73	4.99	4.71	11.69	11.90	10.05	7.9	7.42	7.09				
40	NT2RP2002595	5.43	4.01	5.43	9.33	7.85	7.01	6.61	6.19	7.33	*	+	*	+
	NT2RP2002602	4.82	4.74	4.84	5.43	11.27	8.16	5.69	6.45	7.55			*	+
	NT2RP2002606	5.86	3.02	3.06	8.03	9.33	3.93	3.99	4.72	6.99				
	NT2RP2002609	4.71	2.92	3.43	5.18	4.82	3.59	3.34	4.09	4.4				
	NT2RP2002618	4.82	3.33	2.74	6.13	4.63	4.67	4.95	4.51	4.42				
45	NT2RP2002621	10.26	6.84	5.48	15.22	14.98	13.05	11.07	8.62	10.72	*	+		
	NT2RP2002643	4.22	2.96	3.21	5.73	8.43	4.77	4.53	4.98	4.94			*	+
	NT2RP2002672	4.36	3.45	3.37	8.96	12.04	8.60	8.5	8.50	11.85	**	+	**	+
	NT2RP2002673	2.97	2.38	1.11	7.44	9.35	7.43	5.4	7.46	8.29	**	+	**	+
	NT2RP2002674	1.07	1.16	1.07	0.86	1.66	1.60	1.52	2.10	1.72			*	+
50	NT2RP2002686	3.43	3.39	4.42	4.11	5.80	4.25	4.81	4.16	5.05				
	NT2RP2002688	13.80	10.26	10.39	17.41	16.88	13.34	9.74	11.51	8.03				
	NT2RP2002695	6.80	3.06	3.92	5.81	7.30	4.59	7.03	4.43	5.61				
	NT2RP2002701	6.95	4.89	4.37	8.51	9.98	9.57	8.2	7.53	9.25	*	+	*	+
	NT2RP2002706	4.89	2.72	3.50	5.60	7.16	5.82	4.64	4.20	5.97	*	+		
	NT2RP2002710	42.99	27.04	33.49	47.10	50.09	39.86	54.16	44.21	55.65			*	+
55	NT2RP2002721	7.76	5.23	6.54	9.71	10.64	8.40	10.35	8.03	8.87	*	+		
	NT2RP2002727	0.98	1.45	0.99	2.09	1.15	2.50	2.3	2.04	1.73			*	+

Table 254

	NT2RP2001947	4.91	3.61	5.81	3.96	7.23	5.13	4.97	5.37	4.61			
5	NT2RP2001948	3.08	1.21	4.06	4.99	4.92	1.65	1.37	3.34	8.7			
	NT2RP2001956	15.21	7.64	6.12	7.09	9.06	8.60	13.91	9.28	14.64			
	NT2RP2001969	8.23	4.55	5.29	5.46	6.80	5.70	8.22	5.90	10.07			
	NT2RP2001976	2.14	2.20	2.33	1.64	3.47	2.44	1.48	2.24	2.16			
	NT2RP2001978	4.60	3.86	2.35	6.96	6.45	5.14	6.22	4.96	6.39	*	+	*
	NT2RP2001985	3.92	3.42	3.57	5.93	6.65	5.91	5.3	5.09	5.9	**	+	**
10	NT2RP2001991	1.73	1.46	2.57	3.16	4.44	3.93	3.02	3.02	2.07	*	+	
	NT2RP2001997	3.98	3.95	3.94	5.87	6.12	4.91	4.68	4.05	3.66	*	+	
	NT2RP2002015	78.11	51.57	65.21	141.26	146.10	108.68	76.93	62.92	81.97	**	+	
	NT2RP2002017	3.82	3.00	1.73	4.92	6.18	4.74	4	3.36	3.11	*	+	
	NT2RP2002025	9.38	5.00	3.82	6.47	6.74	7.41	7.27	7.03	6.73			
15	NT2RP2002030	14.24	9.95	8.14	32.58	35.24	33.11	14.46	16.78	20.02	**	+	
	NT2RP2002032	7.60	6.08	6.71	7.52	10.42	7.21	9.78	7.83	10		*	+
	NT2RP2002033	10.00	6.88	8.54	14.32	18.25	17.32	8.01	10.19	9.71	**	+	
	NT2RP2002041	1.30	1.42	1.01	2.33	2.65	2.99	2.24	3.22	3.54	**	+	*
	NT2RP2002046	2.29	2.31	2.63	4.90	5.83	4.05	4.05	4.50	4.31	*	+	*
20	NT2RP2002047	5.55	4.39	6.12	3.39	3.21	2.86	3.07	2.96	1.09	*	-	*
	NT2RP2002050	8.38	3.98	6.12	10.46	10.43	10.14	8.27	8.23	7.23	*	+	
	NT2RP2002052	6.47	4.41	3.60	6.50	9.32	5.86	4.66	4.62	6.58			
	NT2RP2002058	3.62	2.82	3.02	3.46	3.52	2.23	2.78	3.89	2.56			
	NT2RP2002060	6.58	3.14	4.55	4.58	5.81	5.66	5.55	7.36	5.35			
25	NT2RP2002063	1.56	1.90	1.51	3.69	1.67	1.86	2.22	2.63	1.71			
	NT2RP2002066	5.03	3.37	4.61	4.73	5.21	5.32	7.33	6.17	4.62			
	NT2RP2002070	0.79	0.79	0.34	1.28	2.20	1.05	0.97	2.47	0.94			
	NT2RP2002076	3.86	2.57	2.52	3.36	3.56	2.78	2.73	4.09	2.15			
	NT2RP2002078	5.54	3.35	3.42	13.66	10.39	8.08	7.93	6.64	6.4	*	+	*
30	NT2RP2002079	5.14	3.23	1.70	5.80	4.94	6.51	3.67	4.05	3.99			
	NT2RP2002099	7.45	3.48	2.47	4.21	4.13	3.43	3.32	4.93	4.92			
	NT2RP2002105	5.64	3.25	3.05	3.88	4.16	3.68	4.68	5.62	4.37			
	NT2RP2002115	0.92	0.69	0.55	1.83	1.20	1.32	0.97	2.15	0.81	*	+	
	NT2RP2002124	2.28	1.30	1.91	4.70	4.64	3.30	3.98	3.75	2.5	*	+	*
35	NT2RP2002137	2.93	1.88	1.87	2.18	3.16	2.61	3.4	4.11	2.95			
	NT2RP2002139	4.33	3.54	3.42	3.56	4.04	4.02	5.23	4.66	5.13		*	+
	NT2RP2002154	5.53	2.76	1.92	4.83	6.57	3.88	4.83	4.72	5.4			
40	NT2RP2002155	279.79	155.93	163.22	222.28	242.49	184.60	219.6	179.59	177.9			
	NT2RP2002172	4.14	2.59	2.22	3.81	3.52	4.02	3.34	4.90	3.32			
	NT2RP2002185	4.32	3.52	2.95	4.55	4.64	4.41	4.65	5.42	5.45		*	+
	NT2RP2002188	11.41	5.54	8.75	9.54	13.32	9.41	7.96	10.55	9.63			
	NT2RP2002192	3.64	3.48	3.53	4.30	3.68	3.71	1.91	3.83	2.29			
	NT2RP2002193	3.15	2.72	2.77	3.68	4.01	3.41	3.89	3.36	4.16	*	+	*
	NT2RP2002208	2.07	2.36	2.72	6.19	4.41	5.19	4.33	5.08	2.51	**	+	
45	NT2RP2002219	4.17	1.29	1.62	2.78	4.30	2.60	1.31	1.97	1.84			
	NT2RP2002231	2.75	2.39	1.20	3.02	3.57	1.95	2.15	1.47	2.21			
	NT2RP2002232	5.59	1.67	2.23	5.04	5.05	3.28	3.82	4.55	3.16			
	NT2RP2002235	7.15	4.93	3.90	3.84	3.33	2.89	4.86	6.74	5.47			
	NT2RP2002239	23.74	15.37	16.41	23.91	26.96	19.68	8.59	12.98	10.06			
	NT2RP2002252	9.96	4.94	5.61	5.48	5.08	6.39	5.19	6.06	5.88			
50	NT2RP2002256	1.33	1.22	1.37	1.71	2.37	2.14	1.73	2.95	1.47	*	+	
	NT2RP2002257	2.29	1.76	1.74	4.11	5.09	2.83	4.04	4.42	3.81	*	+	**
	NT2RP2002259	3.72	2.30	2.90	6.32	3.45	2.90	3.06	3.48	1.66			
	NT2RP2002264	2.47	1.33	1.14	6.07	7.37	5.74	2.09	3.51	3.03	**	+	
	NT2RP2002267	8.31	4.57	4.68	12.59	14.87	10.14	12.21	9.31	10.07	*	+	*
55	NT2RP2002270	7.39	4.62	5.64	7.88	7.73	8.65	3.38	3.42	4.07			
	NT2RP2002281	8.20	4.58	6.60	7.60	8.32	8.02	5.18	6.11	4.33			
	NT2RP2002288	5.39	5.46	4.44	3.41	3.45	3.50	3.57	3.54	3.96	**	-	-

Table 253

	NT2RP2001560	6.39	4.64	4.20	5.82	7.13	5.81	3.38	4.66	5.13			
	NT2RP2001562	4.89	3.58	3.48	6.44	6.82	4.81	4.71	5.39	5.07			
5	NT2RP2001566	7.48	4.52	5.51	7.16	5.92	8.75	7.73	7.60	6.5			
	NT2RP2001569	14.82	5.79	-9.60	21.83	22.56	14.28	10.25	9.70	10.1			
	NT2RP2001576	10.55	5.49	5.69	8.15	9.33	7.45	8.98	9.68	8.51			
	NT2RP2001581	56.76	28.34	28.83	65.72	65.95	57.58	33.46	29.31	29.57			
	NT2RP2001597	6.52	3.84	3.20	6.75	8.45	4.27	5.43	7.30	6.46			
10	NT2RP2001601	1.39	1.22	0.85	2.84	5.69	3.38	1.83	3.28	2.5	*	+	+
	NT2RP2001613	0.98	1.39	1.71	1.95	1.58	2.25	1.57	2.65	2.69			
	NT2RP2001628	3.83	3.04	3.39	4.74	7.75	4.57	4.66	5.20	3.94			
	NT2RP2001634	9.71	7.65	8.42	9.38	5.92	8.18	7.57	6.78	7.74			
	NT2RP2001635	6.36	3.48	2.24	6.23	7.58	4.38	4.88	3.74	2.85			
15	NT2RP2001660	2.86	2.10	1.03	7.27	5.03	4.32	4.44	3.32	7.02	*	+	
	NT2RP2001662	9.75	5.05	6.57	13.09	11.75	8.88	7.01	6.63	7.59			
	NT2RP2001663	3.29	2.74	2.56	3.86	4.83	6.87	3.87	4.11	4.21		**	+
	NT2RP2001672	3.92	2.66	2.42	6.76	8.23	7.05	3.9	5.21	5.15	**	+	+
	NT2RP2001675	2.35	2.00	2.38	1.25	1.56	1.93	1.59	2.56	2.41	*	-	
20	NT2RP2001677	6.62	5.40	3.75	5.38	8.63	6.75	8.06	7.03	7.46			
	NT2RP2001678	3.81	2.77	2.79	5.76	5.75	5.77	3.78	5.60	5.43	**	+	
	NT2RP2001683	1.31	1.34	1.35	2.92	5.85	2.75	1.53	1.74	1.61		**	+
	NT2RP2001699	10.48	4.46	4.39	9.39	8.26	5.63	7.71	4.72	6.45			
	NT2RP2001707	6.36	2.69	3.12	4.80	5.89	4.38	5.21	3.89	4.02			
25	NT2RP2001720	4.31	2.23	2.64	5.76	5.81	5.36	2.53	3.30	4.19	*	+	
	NT2RP2001721	5.95	3.63	4.33	4.87	4.91	5.43	4.03	4.62	4.71			
	NT2RP2001740	9.64	7.71	6.71	10.42	9.86	6.60	4.64	5.42	6.18			
	NT2RP2001748	8.04	6.16	5.85	6.53	8.57	9.79	7.32	7.38	8.28			
	NT2RP2001755	8.56	5.19	5.01	5.45	6.63	4.59	3	4.11	4.45			
30	NT2RP2001762	3.51	1.45	1.56	4.01	2.49	1.10	1.33	1.59	1.38			
	NT2RP2001768	10.52	5.70	5.26	8.83	8.48	7.75	7.16	7.38	7.69			
	NT2RP2001769	10.19	4.14	4.34	4.02	3.67	3.86	2.04	3.80	3.12			
	NT2RP2001784	3.41	2.66	3.05	4.40	6.83	4.24	3.51	4.60	5.21			
	NT2RP2001805	8.47	4.44	5.36	7.33	9.55	7.18	6.45	7.26	6.85			
35	NT2RP2001813	0.85	0.76	1.30	1.56	0.97	1.22	1.03	2.43	0.53			
	NT2RP2001817	3.31	2.32	3.38	2.20	3.73	2.38	1.83	3.68	1.91			
	NT2RP2001818	9.15	4.97	5.99	7.22	8.04	4.90	5.14	6.97	4.17			
	NT2RP2001837	6.67	3.70	3.89	10.21	8.70	8.64	6.67	5.27	5.41	*	+	
	NT2RP2001839	8.94	4.07	4.05	8.65	8.01	5.90	7.01	4.33	4.71			
40	NT2RP2001861	3.92	3.91	2.96	5.38	4.82	4.41	3.85	3.89	4.28	*	+	
	NT2RP2001869	3.96	3.68	2.84	5.29	6.76	6.36	4.79	4.96	8.38	**	+	
	NT2RP2001876	5.26	4.39	3.67	5.40	6.52	6.44	4.25	3.45	3.89	*	+	
	NT2RP2001878	2.96	2.08	2.84	3.77	3.75	3.70	4.02	3.19	4.69	*	+	
	NT2RP2001881	3.61	3.23	3.04	4.01	3.35	3.50	1.51	1.79	2.14		**	-
45	NT2RP2001883	14.84	8.25	6.92	8.52	8.12	7.84	10.33	7.28	8.44			
	NT2RP2001884	13.60	7.36	6.43	4.80	5.47	5.55	7.44	5.61	6.14			
	NT2RP2001885	4.58	2.98	2.92	4.56	5.26	4.27	4.8	4.09	3.45			
	NT2RP2001898	5.25	3.59	4.61	5.09	5.82	4.63	4.24	6.45	7.13			
	NT2RP2001900	3.76	2.05	3.66	6.01	5.52	2.71	3.58	3.82	6.81			
50	NT2RP2001903	26.27	19.19	22.63	20.41	23.55	21.60	18.49	17.64	17.95			
	NT2RP2001907	6.26	4.16	3.66	9.32	10.90	7.90	6.73	6.46	7.59	*	+	
	NT2RP2001915	2.75	1.61	1.89	3.01	6.15	2.73	2.2	4.12	4.37			
	NT2RP2001921	13.96	7.17	5.50	7.19	5.36	4.44	6.09	4.12	4.96			
	NT2RP2001926	2.31	1.57	1.52	6.10	5.30	3.82	6.1	3.59	5.57	*	+	+
	NT2RP2001933	7.86	5.07	6.52	8.86	5.68	6.54	7.83	5.74	7.63			
55	NT2RP2001936	1.63	0.95	0.99	1.17	2.42	2.36	1.86	2.55	1.83			
	NT2RP2001943	51.19	30.10	31.53	33.01	35.70	30.61	29.07	28.57	30.35			
	NT2RP2001946	3.26	2.65	3.35	3.35	3.83	4.97	4.68	3.30	3.45			

Table 252

	NT2RP2001200	3.43	3.44	2.46	6.55	4.88	4.21	3.59	2.77	3.29			
5	NT2RP2001218	3.11	1.72	2.13	3.51	3.65	3.23	2.31	2.98	3.88			
	NT2RP2001223	5.06	2.55	3.61	3.72	4.59	2.27	3.19	3.20	3.06			
	NT2RP2001226	12.72	7.29	8.85	12.01	9.47	7.65	11.46	8.46	11.8			
	NT2RP2001227	6.22	4.18	3.44	6.26	5.08	5.75	7.03	4.88	5.64			
	NT2RP2001232	7.29	3.90	3.93	7.87	8.17	8.48	7.39	5.90	4.44			
10	NT2RP2001233	14.76	8.17	8.10	14.08	19.00	21.01	13.52	10.12	10.65			
	NT2RP2001245	3.69	2.29	2.63	3.56	3.59	3.28	3.42	3.62	4.39			
	NT2RP2001246	2.35	0.80	3.09	3.34	4.44	4.13	4.38	7.67	6.87		*	+
	NT2RP2001268	5.55	3.73	6.74	8.43	9.77	9.29	5.65	6.17	7.45	*	+	
	NT2RP2001270	14.16	9.13	9.94	14.63	14.49	8.30	11.4	14.47	14.26			
15	NT2RP2001276	2.24	1.82	0.94	3.36	2.75	2.46	3.31	2.32	2.92			
	NT2RP2001277	3.77	1.80	1.15	7.12	6.46	6.90	6.6	4.91	5.92	**	+	*
	NT2RP2001290	3.82	2.12	2.26	5.58	9.49	5.69	6.49	4.65	4.63	*	+	*
	NT2RP2001295	3.75	1.96	2.66	4.93	5.60	3.83	3.62	3.11	3.56			
	NT2RP2001297	104.94	62.95	78.61	112.57	111.95	109.12	28.51	42.30	59.76			
20	NT2RP2001301	6.22	5.96	7.50	7.48	6.39	7.90	5.94	7.38	6.32			
	NT2RP2001312	16.14	10.26	15.91	20.56	19.30	16.72	18.23	19.30	23.86			
	NT2RP2001327	8.14	6.35	5.95	5.76	7.30	7.36	7.73	8.61	9.09			
	NT2RP2001328	18.42	9.64	9.66	24.64	22.08	22.34	13.94	10.86	12.67	*	+	
	NT2RP2001341	17.63	7.30	6.72	12.36	9.62	10.30	8.25	8.97	14.65			
25	NT2RP2001347	17.63	11.15	9.87	16.21	14.33	12.17	10.57	9.73	12.31			
	NT2RP2001346	10.12	8.31	6.45	18.92	23.58	18.36	11.75	11.32	14.59	**	+	*
	NT2RP2001378	8.29	6.95	6.58	6.49	8.22	6.02	7.98	9.16	9.41			
	NT2RP2001381	4.07	2.97	3.94	2.90	3.52	4.42	2.95	2.69	2.85			
	NT2RP2001388	3.41	3.63	3.35	6.25	9.01	7.41	5.95	6.27	6.62	**	+	**
	NT2RP2001391	210.40	161.64	144.04	393.09	492.35	288.04	175.7	224.46	230.6	*	+	
30	NT2RP2001392	7.04	3.01	3.58	4.59	5.33	4.71	6.14	5.70	5.27			
	NT2RP2001394	9.60	6.22	4.32	15.24	15.30	14.78	8	5.76	7.4	**	+	
	NT2RP2001397	15.57	11.63	10.83	8.23	11.47	9.12	4.18	3.62	3.82		**	-
	NT2RP2001400	2.42	2.39	2.33	4.87	6.19	6.06	7.4	8.87	13.18	**	+	*
	NT2RP2001408	5.20	3.88	3.54	7.39	10.57	7.94	7.53	7.30	6.48	*	+	**
35	NT2RP2001420	4.15	2.99	3.26	8.92	7.75	7.19	4.98	4.32	3.55	**	+	
	NT2RP2001423	3.65	2.45	3.55	6.47	6.38	4.42	6.23	5.04	5.49	*	+	*
	NT2RP2001427	4.90	3.28	3.58	5.81	6.42	5.73	4.13	4.89	4.51	*	+	
	NT2RP2001428	4.31	2.09	2.32	7.25	7.90	5.77	3.53	5.08	3.14	*	+	
	NT2RP2001436	3.76	2.25	2.26	8.78	8.61	8.75	5.22	4.80	6.42	**	+	*
40	NT2RP2001440	3.29	2.41	1.73	3.63	4.88	4.33	2.34	3.35	3.86	*	+	
	NT2RP2001445	2.95	1.26	2.68	2.98	3.78	3.07	2.47	3.15	2.23			
	NT2RP2001449	2.88	2.13	1.40	3.15	3.39	4.62	2.6	3.60	1.97			
	NT2RP2001450	4.05	2.94	3.13	3.77	4.91	3.85	3.71	4.15	3.13			
	NT2RP2001467	2.37	1.91	2.75	5.44	4.55	6.16	5.15	4.88	3.4	**	+	*
45	NT2RP2001469	10.04	7.34	9.26	5.41	8.75	6.36	6.52	6.42	6.37		*	-
	NT2RP2001480	6.23	4.15	2.86	6.30	5.94	4.07	6.36	5.86	4.85			
	NT2RP2001495	14.26	10.91	10.35	11.90	13.38	11.11	12.39	11.10	12.13			
	NT2RP2001499	4.67	3.29	2.95	6.59	7.16	8.76	5.49	6.33	5.02	**	+	*
	NT2RP2001506	4.89	3.71	3.86	7.29	8.04	7.88	5.96	6.72	7.88	**	+	*
	NT2RP2001508	6.85	6.36	6.72	17.18	14.22	13.59	7.65	11.84	6.81	**	+	
50	NT2RP2001511	11.59	6.20	8.17	12.86	12.37	12.22	11.15	9.45	10.6			
	NT2RP2001514	6.61	4.54	5.10	6.50	5.89	6.49	5.87	6.17	7.22			
	NT2RP2001520	2.37	1.99	2.43	3.12	2.75	2.57	4.35	3.09	2.57			
	NT2RP2001526	12.96	5.00	5.99	26.60	29.55	19.41	14.77	8.09	13.41	*	+	
	NT2RP2001529	8.76	6.16	5.20	6.03	7.62	5.55	4.77	5.25	10.17			
55	NT2RP2001536	3.16	2.19	1.50	3.33	3.18	2.23	2.35	2.79	2.24			
	NT2RP2001538	75.84	48.30	57.88	103.08	97.23	96.04	48.73	42.09	45.9	*	+	
	NT2RP2001547	5.37	2.86	3.64	4.76	4.73	4.52	3.5	4.96	4.64			

Table 251

	NT2RP2000777	29.85	20.42	21.91	16.22	17.42	15.02	12.37	12.07	13.5		*	-
5	NT2RP2000786	8.23	5.22	4.46	10.55	9.74	7.80	11.68	12.09	10.59		*	+
	NT2RP2000793	14.01	7.42	10.26	12.19	18.41	17.74	12.21	12.81	15.75			
	NT2RP2000796	6.25	2.57	4.14	5.05	5.14	3.86	3.27	4.71	3.04			
	NT2RP2000809	7.70	5.02	4.14	9.32	10.55	8.44	6.87	4.85	6.03	*	+	
	NT2RP2000812	6.41	3.65	3.75	7.05	6.14	5.86	4.83	5.04	3.67			
10	NT2RP2000814	2.40	1.13	1.50	2.03	1.96	1.78	0.9	2.54	1.09			
	NT2RP2000816	5.89	1.17	2.01	3.48	3.06	4.82	3.84	4.28	3.29			
	NT2RP2000818	2.61	0.71	0.86	3.13	3.87	2.75	2.08	1.63	3.25			
	NT2RP2000819	2.57	1.24	1.34	1.88	1.49	1.77	1.81	2.05	1.32			
	NT2RP2000841	2.46	0.72	1.21	2.94	1.98	3.02	1.06	2.75	1.48			
15	NT2RP2000842	1.34	0.54	1.09	1.95	1.45	1.71	2.84	2.70	1.53		*	+
	NT2RP2000845	12.78	5.61	3.57	11.56	12.23	11.13	7.34	7.10	8.72			
	NT2RP2000843	2.24	1.48	1.52	2.02	1.72	1.96	1.61	2.25	1.68			
	NT2RP2000880	10.87	4.76	7.03	10.28	10.84	10.60	7.87	8.04	7.97			
	NT2RP2000892	3.07	1.45	2.10	2.15	3.52	2.03	2.6	3.34	2.68			
20	NT2RP2000894	2.45	1.27	1.87	2.80	3.03	2.60	3.77	5.13	5.17		**	+
	NT2RP2000903	2.42	1.74	2.17	15.91	10.43	12.06	3.76	4.80	3.91	**	+	**
	NT2RP2000906	2.89	1.95	2.70	4.14	5.17	4.16	3.32	2.67	4.12	*	+	
	NT2RP2000910	2.79	1.53	2.66	6.17	5.30	4.67	3.71	4.07	3.28	**	+	*
	NT2RP2000931	32.13	11.92	13.53	39.97	39.93	28.59	17.58	15.27	16.3			
	NT2RP2000932	4.21	2.31	2.05	7.96	6.87	4.87	4.36	3.76	4.67	*	+	
25	NT2RP2000938	19.54	10.59	13.57	13.71	16.06	13.76	9.46	10.81	12.03			
	NT2RP2000943	4.61	2.00	2.25	2.99	4.17	3.48	6.66	6.59	6.2		*	+
	NT2RP2000957	2.25	1.38	1.92	2.45	2.33	2.46	1.28	3.48	2.23			
	NT2RP2000958	6.62	2.75	4.11	5.71	4.71	5.65	4.44	6.65	3.45			
30	NT2RP2000959	5.43	1.74	2.79	6.81	7.31	5.96	7.7	6.58	8.28	*	+	*
	NT2RP2000965	8.62	7.11	7.91	6.90	6.39	7.29	4.61	4.19	4.83		**	-
	NT2RP2000970	6.70	2.82	2.67	8.85	8.32	8.60	5.68	4.48	4.57	*	+	
	NT2RP2000973	3.87	3.35	2.21	3.68	3.61	1.94	3.33	3.24	2.43			
	NT2RP2000985	4.15	2.39	2.33	2.87	4.28	3.35	2.71	2.53	3.95			
	NT2RP2000987	2.36	1.40	1.29	2.94	3.30	3.87	2.43	3.02	3.28	*	+	*
35	NT2RP2000997	3.92	3.46	2.91	6.76	6.13	8.29	6.06	7.63	6.82	**	+	**
	NT2RP2001024	3.02	2.00	2.80	4.39	4.00	3.80	2.57	2.72	3.03	*	+	
	NT2RP2001028	1.53	1.61	1.49	3.31	2.89	2.16	1.09	3.10	1.56	*	+	
	NT2RP2001036	8.99	5.09	6.28	14.47	12.09	13.66	6.21	7.37	8.86	**	+	
	NT2RP2001039	2.38	1.24	0.84	2.83	2.64	1.64	1.85	1.41	1.82			
40	NT2RP2001044	3.60	1.75	2.33	3.81	3.95	2.60	1.92	3.42	3.51			
	NT2RP2001056	8.76	6.20	3.80	10.38	10.96	8.29	5.85	5.19	6.9			
	NT2RP2001065	11.06	6.53	6.66	6.07	7.52	5.67	4.84	4.18	3.98			
	NT2RP2001067	3.97	2.56	1.95	4.29	2.72	3.44	1.28	3.38	2.55			
	NT2RP2001070	6.27	3.18	2.94	8.92	8.75	6.08	5.11	6.42	3.18			
	NT2RP2001081	7.29	3.39	2.85	9.20	10.42	10.02	6.26	8.11	6.41	*	+	
45	NT2RP2001087	2.47	2.17	1.24	3.46	5.06	3.87	2.98	3.13	3.05	*	+	*
	NT2RP2001094	0.61	0.13	0.10	1.14	0.70	0.35	0.83	0.86	1.21		*	+
	NT2RP2001119	6.84	4.46	3.47	7.70	9.69	7.83	4.19	5.13	8.84	*	+	
	NT2RP2001127	5.97	3.17	2.14	8.14	7.01	6.94	3.37	5.51	5.47	*	+	
	NT2RP2001133	6.80	4.14	3.76	7.22	8.84	6.01	3.82	6.62	4.59			
50	NT2RP2001137	4.85	2.38	2.65	2.75	3.98	3.93	2.74	5.27	3.23			
	NT2RP2001142	3.86	1.91	2.02	3.11	3.09	2.46	1.97	4.83	1.3			
	NT2RP2001149	4.02	1.34	2.11	3.88	2.95	3.29	1.85	2.88	2.53			
	NT2RP2001168	13.95	5.65	7.80	16.05	15.12	13.54	11.11	11.37	10.13			
	NT2RP2001173	2.96	1.32	1.35	7.72	6.56	4.53	4.19	3.26	2.72	*	+	
55	NT2RP2001174	4.49	3.17	1.74	5.69	5.38	5.09	5.65	3.56	3.21			
	NT2RP2001184	7.71	4.21	4.96	7.15	6.32	5.98	5.09	5.61	5.63			
	NT2RP2001196	1.68	0.99	1.05	1.56	1.51	1.49	1.6	1.79	2.14			

Table 250

	NT2RP2000422	4.34	2.42	2.61	4.23	4.79	3.97	2.48	4.12	3.58			
5	NT2RP2000426	25.72	16.73	17.55	38.01	37.89	27.90	28.44	35.63	32.72	*	+	+
	NT2RP2000428	8.81	5.15	7.26	4.95	7.26	4.98	5.88	6.67	7.85			
	NT2RP2000438	6.31	4.25	6.08	7.20	6.52	5.26	4.94	5.80	4.64			
	NT2RP2000447	4.41	2.06	2.07	4.91	3.95	2.02	2.15	2.90	4.07			
	NT2RP2000448	7.83	4.29	4.32	8.83	10.57	6.61	6.83	6.72	9.81			
	NT2RP2000459	3.66	2.01	1.92	4.90	4.18	3.40	3.04	3.12	2.39			
10	NT2RP2000479	1.93	0.77	1.02	3.37	3.48	3.07	1.64	3.13	2.2	**	+	
	NT2RP2000498	3.73	1.64	2.79	6.08	6.58	5.26	3.06	4.66	3.3	*	+	
	NT2RP2000503	0.99	0.59	0.90	1.83	1.74	0.79	1.01	2.91	0.59			
	NT2RP2000510	1.06	0.59	0.92	1.09	1.85	1.43	0.94	2.45	1.3			
	NT2RP2000514	1.41	1.10	1.00	1.62	1.02	0.66	0.8	2.20	1.21			
15	NT2RP2000516	2.96	2.89	1.64	2.85	2.86	3.71	3.31	2.30	4			
	NT2RP2000523	3.99	1.92	2.37	1.57	3.10	1.25	3.93	1.39	1.65			
	NT2RP2000533	8.58	5.78	6.04	9.66	6.29	7.70	8.46	6.89	6.1			
	NT2RP2000540	3.70	1.50	1.36	1.88	3.29	2.35	3	2.34	2.25			
	NT2RP2000547	4.21	3.25	2.00	3.94	5.17	3.32	3.43	3.90	3.44			
20	NT2RP2000557	6.17	3.16	5.21	9.43	7.58	8.00	4.94	5.68	5.75	*	+	
	NT2RP2000558	6.82	5.39	2.81	8.42	7.99	7.74	3.91	5.66	3.66			
	NT2RP2000564	3.37	1.73	2.60	5.24	4.86	4.91	2.08	2.76	4.62	**	+	
	NT2RP2000565	10.89	3.85	5.45	5.34	4.15	3.62	5.93	5.18	4.1			
	NT2RP2000583	12.11	7.48	7.41	14.37	9.94	10.68	9.35	8.42	9.2			
25	NT2RP2000591	1.21	1.15	0.59	1.83	2.04	1.49	1.94	1.98	1.05	*	+	
	NT2RP2000599	1.47	1.25	1.53	1.16	1.55	1.34	1.22	2.03	0.81			
	NT2RP2000601	2.53	1.94	2.56	4.22	3.80	2.72	5.23	4.02	4.33		**	+
	NT2RP2000603	3.39	2.35	1.65	2.95	3.86	3.73	3.27	3.61	3.79			
	NT2RP2000610	8.35	6.25	7.50	11.79	10.08	10.19	6.69	6.74	5.04	*	+	
30	NT2RP2000614	96.26	103.19	118.68	120.08	119.37	64.42	36.46	62.71	38.98		**	-
	NT2RP2000616	6.76	3.07	4.14	4.68	4.17	3.26	5.28	4.32	4.63			
	NT2RP2000617	8.33	3.91	4.08	4.27	5.55	4.60	5.01	3.15	4.64			
	NT2RP2000623	4.48	1.59	1.85	3.07	2.65	2.79	2.55	2.58	1.9			
	NT2RP2000634	2.21	1.66	0.95	4.67	6.41	3.91	3.28	3.56	3.18	*	+	+
35	NT2RP2000636	2.78	1.86	2.23	5.39	5.75	3.65	5.59	4.74	6.43	*	+	**
	NT2RP2000638	21.16	12.92	16.03	4.08	3.49	3.77	3.77	2.86	3.58	**	-	**
	NT2RP2000644	4.37	1.59	2.30	6.98	6.00	7.24	4.21	4.56	3.58	*	+	
	NT2RP2000649	7.14	4.82	5.18	7.37	7.32	4.24	9.38	7.32	6.55			
	NT2RP2000652	3.51	2.62	3.37	2.59	3.37	3.58	3.42	2.20	3.62			
	NT2RP2000656	2.66	3.06	2.65	4.78	6.50	7.33	2.65	3.45	3.99	*	+	
40	NT2RP2000658	0.93	1.13	0.36	1.13	1.33	1.51	1.68	1.25	0.75			
	NT2RP2000663	4.22	2.97	3.08	9.06	10.89	6.58	6.13	6.43	9.35	*	+	+
	NT2RP2000664	23.91	17.42	14.73	9.66	12.53	10.44	7.05	5.83	8.31		*	-
	NT2RP2000668	5.30	2.81	4.65	6.71	5.59	4.69	6.21	4.52	4.52			
	NT2RP2000678	0.48	0.48	0.42	0.75	0.94	0.64	0.81	1.41	0.39	*	+	
45	NT2RP2000694	2.29	2.24	2.05	19.86	17.58	12.78	4.53	4.69	3.6	**	+	**
	NT2RP2000704	6.91	3.49	2.43	6.07	5.63	5.83	4.96	5.30	4.17			
	NT2RP2000710	9.01	4.65	4.93	4.63	5.99	4.41	2.4	3.05	3.57			
	NT2RP2000712	8.69	3.86	3.32	7.90	11.98	9.71	4.72	4.64	4.82			
	NT2RP2000715	2.82	2.17	1.75	4.86	5.63	4.47	3.49	4.30	2.59	**	+	
50	NT2RP2000720	4.75	3.62	3.91	5.03	5.10	4.87	4.06	4.30	4.01			
	NT2RP2000731	2.07	0.87	1.19	1.70	1.63	1.57	2.2	2.35	1.11			
	NT2RP2000739	4.43	3.04	3.85	4.97	5.04	12.49	4.91	4.00	4.7			
	NT2RP2000748	2.01	0.84	1.62	3.67	2.92	3.60	1.71	2.82	2.4	*	+	
	NT2RP2000749	18.07	9.30	9.03	17.51	22.01	17.32	13.66	13.69	16.17			
55	NT2RP2000758	6.82	2.65	3.39	7.55	7.28	7.11	5.45	4.30	5.82			
	NT2RP2000764	6.06	3.40	3.08	3.96	3.81	2.89	3.82	5.32	3.35			
	NT2RP2000766	4.46	2.57	3.04	28.36	19.71	19.14	14.72	13.56	10.75	**	+	**

Table 249

	NT2RP2000119	8.68	3.95	4.21	9.78	9.83	7.70	4.38	5.61	4.76			
	NT2RP2000120	6.77	5.63	5.88	9.79	11.11	8.08	7.54	6.05	5.79 *	+		
5	NT2RP2000126	6.86	4.89	4.70	8.53	5.94	6.57	4.76	5.23	4.11			
	NT2RP2000133	3.99	1.70	2.52	3.67	4.08	3.28	3.34	3.20	1.96			
	NT2RP2000147	10.14	5.06	4.39	7.57	6.45	7.93	7.96	5.91	7.47			
	NT2RP2000153	9.59	4.30	4.77	11.17	12.10	9.91	6.51	6.58	8.83			
	NT2RP2000156	8.43	4.96	3.48	10.08	10.36	9.94	5.38	4.40	3.72 *	+		
10	NT2RP2000157	3.42	2.19	2.41	3.80	5.30	4.72	2.87	2.06	2.91 *	+		
	NT2RP2000161	3.63	2.23	2.07	2.95	5.95	3.11	2.97	3.99	3.8			
	NT2RP2000168	0.99	0.64	1.00	1.63	1.21	0.85	1.57	2.63	1.12			
	NT2RP2000173	5.26	3.38	4.83	5.31	6.20	4.30	6.86	7.09	4.77			
	NT2RP2000175	5.66	3.98	5.08	6.59	5.28	4.03	5.09	5.43	4.57			
15	NT2RP2000178	4.05	2.68	1.96	2.97	4.24	3.15	4.17	4.26	3.99			
	NT2RP2000183	10.17	3.83	4.48	9.26	9.55	10.17	7.2	6.57	6.26			
	NT2RP2000195	7.49	2.50	2.99	9.64	9.13	9.97	5.54	5.28	4.35 *	+		
	NT2RP2000204	61.75	38.58	41.68	97.90	112.72	86.99	46.74	43.39	38.72 **	+		
	NT2RP2000205	3.47	1.89	2.20	5.10	3.54	4.32	2.79	2.79	2.7			
20	NT2RP2000208	3.13	2.58	1.85	5.38	5.41	5.54	3.65	4.43	4.57 **	+	*	+
	NT2RP2000224	10.06	4.94	5.26	13.62	13.47	11.09	7.3	8.43	8.25 *	+		
	NT2RP2000230	10.44	5.32	7.82	4.62	4.88	4.53	6.76	7.92	6.25			
	NT2RP2000231	15.70	8.92	8.46	8.81	11.88	10.86	12.38	9.81	14.32			
	NT2RP2000232	3.82	2.08	1.56	2.18	2.93	2.14	2.17	3.16	3.23			
25	NT2RP2000233	3.92	2.50	2.55	3.87	3.62	3.14	4.2	5.00	3.42			
	NT2RP2000239	5.63	2.55	4.01	2.51	2.65	1.68	2.58	2.65	2.15			
	NT2RP2000240	2.65	0.99	1.49	3.74	2.57	2.17	1.29	3.46	1.94			
	NT2RP2000248	2.07	1.21	1.92	5.23	4.26	2.91	2.54	3.82	2.58 *	+		
	NT2RP2000256	2.45	1.19	2.67	4.07	3.99	4.15	2.35	4.00	2.51 *	+		
30	NT2RP2000257	4.01	2.58	4.00	7.82	7.06	6.67	4.5	7.31	5.28 **	+		
	NT2RP2000258	4.50	2.39	2.97	2.52	3.60	4.01	2.36	1.90	2.05			
	NT2RP2000261	5.05	1.91	1.66	2.79	3.32	2.35	3.34	3.46	3.43			
	NT2RP2000270	4.76	3.28	4.00	7.87	7.75	6.15	4.27	5.23	5.14 **	+		
	NT2RP2000274	1.79	1.60	1.36	2.19	2.83	2.80	2.75	3.55	2.34 *	+	*	+
	NT2RP2000277	2.75	1.21	1.42	2.17	1.68	1.96	1.92	2.84	2.38			
35	NT2RP2000279	0.41	1.31	1.45	1.18	1.47	1.06	1.2	2.43	1.11			
	NT2RP2000283	3.37	2.23	2.52	5.72	4.12	4.64	3.18	4.04	2.42 *	+		
	NT2RP2000288	5.70	4.02	4.20	8.50	6.14	8.35	4.51	3.57	3.55 *	+		
	NT2RP2000289	6.80	5.85	3.10	6.12	5.47	3.78	3.88	3.57	4.36			
	NT2RP2000297	11.76	5.46	4.79	20.39	23.99	16.10	8.54	7.85	6.17 *	+		
40	NT2RP2000298	4.88	2.68	4.30	8.97	6.69	7.77	3.27	4.79	4.52 *	+		
	NT2RP2000310	3.32	1.70	1.94	1.61	2.82	2.27	1.42	3.61	2.47			
	NT2RP2000327	2.70	2.09	1.98	2.16	2.54	2.15	1.73	3.66	2.67			
	NT2RP2000328	9.99	5.11	5.84	9.30	7.53	6.17	5.88	5.38	4.93			
	NT2RP2000329	6.52	3.59	6.38	14.80	8.75	11.24	11.8	13.63	15.25 *	+	**	+
45	NT2RP2000333	2.61	2.37	2.88	3.29	2.69	3.44	2.94	4.19	2.52			
	NT2RP2000337	1.84	1.24	0.70	1.53	2.14	1.62	1.08	1.19	1.29			
	NT2RP2000346	6.13	3.16	4.39	6.09	6.33	4.39	5.29	3.87	4.75			
	NT2RP2000357	4.83	1.57	2.53	4.81	4.10	3.76	2.25	2.94	2.98			
	NT2RP2000358	4.05	2.01	1.43	3.71	5.44	4.47	2.33	3.23	3.82			
50	NT2RP2000366	3.62	3.12	2.58	3.24	4.65	4.15	3.46	5.12	4			
	NT2RP2000369	3.68	3.14	3.25	7.30	6.97	6.80	16.68	15.91	21.03 **	+	**	+
	NT2RP2000376	16.50	7.18	10.26	12.72	14.14	12.56	11.16	13.27	14.04			
	NT2RP2000394	3.97	3.08	4.07	2.94	3.29	3.97	2.41	3.13	3.01			
	NT2RP2000396	14.08	6.54	5.86	11.48	9.74	7.82	9.11	5.57	11.18			
55	NT2RP2000412	7.77	4.65	2.97	6.62	7.16	4.26	3.14	4.29	4.91			
	NT2RP2000414	18.85	9.88	9.70	17.32	11.45	11.38	9.42	7.23	10.75			
	NT2RP2000420	2.85	2.26	2.25	4.04	3.82	1.85	2.03	3.71	2.9			

Table 248

	NT2RP1001424	2.87	1.62	0.72	3.11	2.58	2.58	1.61	3.38	2.21				
	NT2RP1001432	2.47	1.17	2.41	2.23	2.48	1.53	1.78	3.14	1.45				
5	NT2RP1001449	7.62	4.22	5.10	9.69	11.61	8.75	6.99	5.82	6.74	*	+		
	NT2RP1001457	4.04	2.37	2.71	3.08	3.14	2.75	2.72	2.61	3.14				
	NT2RP1001459	10.76	3.49	3.82	8.95	9.17	5.61	7.87	6.73	6.96				
	NT2RP1001466	22.82	9.71	11.08	9.67	7.98	7.40	7.72	5.26	6.18				
	NT2RP1001475	6.67	4.07	4.28	8.53	10.26	8.11	4.73	4.80	4.35	*	+		
10	NT2RP1001482	11.57	4.98	6.24	6.89	5.62	4.62	2.44	2.41	2.61				
	NT2RP1001494	1.38	1.05	0.95	2.03	1.52	1.37	0.9	2.19	2.18				
	NT2RP1001500	2.19	2.12	1.80	1.11	1.95	1.39	1	2.88	1.81				
	NT2RP1001517	1.81	0.96	1.45	2.37	1.81	2.59	1.22	2.90	1.19				
	NT2RP1001540	5.66	2.57	3.71	5.28	5.66	5.56	4.29	5.21	3.47				
15	NT2RP1001543	8.78	3.57	3.35	10.80	11.59	6.71	5.01	4.67	5.34				
	NT2RP1001546	21.79	10.60	8.72	53.53	51.38	41.78	29.72	27.77	37.59	**	+	*	+
	NT2RP1001550	9.54	5.59	4.56	11.19	14.45	11.40	5.56	6.79	7.13	*	+		
	NT2RP1001553	6.39	3.38	2.69	4.45	3.49	2.74	3.6	4.16	2.78				
	NT2RP1001555	9.92	5.57	6.23	12.43	10.74	10.45	11.69	18.08	11.64				
20	NT2RP1001563	4.37	1.97	2.43	3.66	4.03	3.10	1.84	4.08	2.22				
	NT2RP1001569	5.25	3.17	2.27	4.32	4.47	4.21	3.54	6.70	3.89				
	NT2RP1001584	8.28	4.33	4.71	6.70	8.09	6.25	5.94	7.17	6.75				
	NT2RP1001599	7.22	2.05	1.29	32.60	27.43	19.18	6.56	7.36	8.71	**	+		
	NT2RP1001616	3.29	0.83	1.26	2.03	2.10	1.09	2.49	2.44	3.45				
25	NT2RP1001654	19.86	5.14	4.62	10.80	10.51	8.45	6.66	9.40	9.83				
	NT2RP1001665	1.29	1.28	0.35	1.08	1.87	1.90	0.74	2.81	0.7				
	NT2RP1001679	87.88	43.02	42.15	72.20	73.59	55.81	28.48	41.49	35.04				
	NT2RP1001681	21.69	14.86	18.60	13.78	17.18	10.98	7.11	14.27	11.13				
	NT2RP1001694	8.51	6.03	4.96	4.21	4.41	2.94	5.31	11.65	6.79				
30	NT2RP2000001	6.32	1.40	2.79	3.24	2.80	2.62	3.54	4.14	4.08				
	NT2RP2000006	2.04	1.48	0.96	4.50	2.92	2.33	2.69	2.50	1.61				
	NT2RP2000007	10.09	4.44	5.04	3.97	3.31	4.03	3.55	1.69	1.81				
	NT2RP2000008	10.88	5.03	5.27	12.65	14.30	9.35	7.5	5.73	4.32				
	NT2RP2000010	1.99	1.02	0.52	2.09	3.06	2.49	2.1	2.41	2.6				
35	NT2RP2000011	7.02	4.29	5.02	10.56	10.46	8.08	6.55	5.23	6.43	*	+		
	NT2RP2000027	3.12	1.86	1.41	5.78	3.32	2.95	2.99	2.41	1.39				
	NT2RP2000028	2.89	1.81	1.90	3.51	2.63	2.74	4.34	5.32	5.48		**	+	
	NT2RP2000032	1.94	1.20	2.03	2.85	3.78	4.04	1.05	2.10	0.96	*	+		
	NT2RP2000040	37.68	15.23	16.54	19.89	18.06	16.95	22.42	19.65	16.11				
	NT2RP2000042	9.28	3.40	4.33	7.54	7.04	6.30	5.89	6.48	6.12				
40	NT2RP2000045	10.41	4.33	5.29	6.44	6.23	7.07	5.45	5.93	4.14				
	NT2RP2000051	12.68	6.63	7.07	5.35	6.94	5.58	5.26	5.53	4.86				
	NT2RP2000054	5.27	3.29	2.87	3.98	5.04	4.42	5.28	3.65	4.48				
	NT2RP2000056	4.49	2.47	2.46	3.36	3.01	3.82	3.5	3.62	3.48				
	NT2RP2000057	52.52	38.64	47.28	59.49	56.29	50.39	23.72	29.59	31.7		*	.	
45	NT2RP2000067	3.42	1.83	2.49	4.64	3.08	3.41	1.5	3.38	2.02				
	NT2RP2000070	8.99	4.22	3.23	5.71	5.95	7.00	8.23	3.07	7.09				
	NT2RP2000076	2.83	1.15	1.15	1.86	1.61	1.73	2.7	1.97	2.2				
	NT2RP2000077	10.69	4.72	3.55	9.58	8.73	8.11	7.3	4.40	8.28				
	NT2RP2000079	4.88	3.21	3.11	8.07	7.12	7.59	4.5	3.56	4.48	**	+		
50	NT2RP2000083	3.87	3.74	2.96	4.10	4.22	2.91	4.51	4.30	4.17				
	NT2RP2000091	3.05	2.14	3.45	10.95	9.06	8.83	4.37	6.05	6.43	**	+	*	+
	NT2RP2000092	10.83	5.23	7.63	16.92	17.59	12.32	8.03	11.12	9.14	*	+		
	NT2RP2000097	2.33	2.76	2.63	4.90	4.82	3.90	2.22	2.43	2.99	**	+		
	NT2RP2000098	10.38	5.79	6.50	5.56	4.26	4.65	2.67	1.61	2.03		*	.	
55	NT2RP2000108	9.83	5.39	6.38	12.17	15.62	9.37	8.01	6.04	4.82				
	NT2RP2000114	2.05	1.50	1.13	3.20	1.92	2.20	3.45	2.13	2.56				
	NT2RP2000116	5.05	3.16	5.23	7.97	9.36	8.63	7.01	7.36	8.27	**	+	*	+

Table 247

	NT2RP1000916	3.20	2.97	0.92	3.11	2.12	2.20	3.16	2.17	2.19			
	NT2RP1000921	1.84	1.45	1.78	3.53	2.23	2.78	2.9	2.99	2.53	*	+	**+
5	NT2RP1000943	1.83	0.78	1.29	5.94	5.07	4.31	7.05	7.60	6.55	**	+	**+
	NT2RP1000944	3.54	2.52	3.09	5.21	4.55	4.65	2.55	2.42	2.55	**	+	
	NT2RP1000947	6.99	4.11	3.31	6.97	6.41	5.03	5.81	4.15	4.54			
	NT2RP1000954	5.12	2.35	2.15	5.93	4.95	4.84	4.75	3.63	4.18			
	NT2RP1000958	20.62	10.44	1.43	11.21	10.24	6.49	7.05	5.48	7.18			
10	NT2RP1000959	72.56	35.16	43.30	53.44	48.85	40.35	20.64	19.16	22.61			
	NT2RP1000966	36.86	19.10	21.19	22.56	35.39	24.14	15.07	9.91	18.23			
	NT2RP1000974	10.91	8.14	8.28	18.92	22.10	19.21	14.69	15.24	13.39	**	+	**+
	NT2RP1000980	3.63	2.59	2.91	3.75	4.02	3.96	2.97	3.22	2.22			
	NT2RP1000981	4.96	3.42	4.61	4.59	5.02	3.62	2.94	3.11	2.77		*	-
15	NT2RP1000988	2.69	1.97	1.73	4.25	5.22	4.19	3.95	3.30	3.66	**	+	*+
	NT2RP1001002	6.75	4.73	2.89	3.13	4.46	2.79	4.86	5.58	5.21			
	NT2RP1001004	1.76	1.26	0.75	1.72	1.80	2.22	3.2	2.14	2.89		*	+
	NT2RP1001007	1.72	0.91	0.86	2.02	1.84	1.75	3	2.58	3.22		**	+
	NT2RP1001011	4.98	3.03	2.17	7.06	8.67	6.46	5.23	4.65	5.76	*	+	
20	NT2RP1001013	3.60	3.50	3.48	9.46	12.09	7.99	6.88	5.63	8.02	**	+	**+
	NT2RP1001014	3.96	3.16	3.28	4.93	3.71	4.01	3.71	3.05	2.43			
	NT2RP1001020	3.23	1.24	1.06	2.23	1.86	1.47	2.29	2.09	1.68			
	NT2RP1001023	261.06	118.84	124.95	113.92	104.93	83.66	236.2	219.46	213.5			
	NT2RP1001027	12.10	6.08	4.74	9.03	7.91	6.47	4.01	4.15	4.04			
25	NT2RP1001031	2.17	1.05	0.67	1.79	1.31	1.73	0.62	1.86	1.33			
	NT2RP1001033	2.89	1.62	1.96	3.31	4.49	3.57	2.4	3.46	2.46	*	+	
	NT2RP1001042	2.56	1.34	2.04	5.44	5.57	4.27	4.68	4.47	4.7	**	+	**+
	NT2RP1001045	55.87	37.46	39.12	31.66	32.21	26.52	26.73	25.41	27.84		*	-
	NT2RP1001073	18.17	10.94	13.65	7.43	11.62	10.45	3.3	5.82	3.6		*	-
	NT2RP1001079	6.27	4.29	4.83	7.17	5.68	5.81	5.84	5.09	3.64			
30	NT2RP1001080	4.59	3.36	2.02	3.32	2.67	3.66	3.81	3.01	2.62			
	NT2RP1001113	2.09	1.06	0.43	0.85	1.89	1.25	1.74	2.63	1.22			
	NT2RP1001159	22.23	15.34	13.51	27.36	29.04	20.75	11.14	12.23	9.12			
	NT2RP1001173	2.37	0.91	1.48	10.20	7.72	8.04	6.93	5.00	6.33	**	+	**+
35	NT2RP1001176	5.14	3.86	5.35	6.46	6.12	5.31	4.46	5.39	4.12			
	NT2RP1001177	3.79	2.64	3.45	7.23	6.84	5.24	5.18	4.11	3.16	*	+	
	NT2RP1001185	4.77	2.20	2.83	10.28	7.74	6.42	4.72	4.39	3.75	*	+	
	NT2RP1001199	2.06	1.25	1.14	4.62	4.88	3.76	2.05	2.71	1.7	**	+	
	NT2RP1001205	19.37	11.82	11.58	17.19	17.16	12.69	6.66	6.05	4.62		*	-
	NT2RP1001215	5.66	2.61	2.14	2.79	3.86	3.71	2.65	3.10	2.8			
40	NT2RP1001225	5.42	2.06	1.65	2.88	2.39	2.40	3.21	4.49	4.21			
	NT2RP1001245	3.12	2.43	4.04	4.32	4.51	4.91	3.1	5.42	4.42	*	+	
	NT2RP1001247	1.41	0.44	0.55	0.62	0.90	1.10	0.75	2.81	1			
	NT2RP1001248	2.68	2.07	1.62	3.98	2.41	2.41	1.39	3.80	1.81			
	NT2RP1001253	6.69	3.25	3.71	6.33	4.35	5.83	4.57	5.25	3.74			
45	NT2RP1001286	3.18	1.26	2.31	4.52	3.67	4.87	3.96	3.81	2.61	*	+	
	NT2RP1001294	9.78	2.41	4.54	3.50	4.67	2.27	2.68	1.87	2.59			
	NT2RP1001302	8.57	3.22	3.02	3.18	3.24	3.37	2.74	2.23	2.95			
	NT2RP1001310	9.73	5.23	5.10	9.63	10.00	7.15	7.46	7.70	6.61			
	NT2RP1001311	18.47	7.91	7.87	5.75	8.43	7.25	3.98	5.42	3.54			
50	NT2RP1001313	10.94	5.16	4.72	12.65	11.32	9.22	3.47	5.55	4.61			
	NT2RP1001324	3.38	2.26	1.54	3.44	2.03	2.97	2.03	3.34	1.99			
	NT2RP1001349	3.51	1.77	2.13	2.29	2.35	2.91	2.76	4.09	1.76			
	NT2RP1001361	9.53	5.57	12.07	15.75	14.43	10.15	3.96	7.68	5.72			
	NT2RP1001379	9.49	3.63	4.16	6.43	5.54	3.66	4.65	4.16	4.18			
	NT2RP1001385	6.18	2.32	2.60	4.81	6.35	3.73	2.76	3.62	3.67			
55	NT2RP1001395	5.45	2.82	3.04	4.04	3.63	2.71	4.99	4.64	3.44			
	NT2RP1001410	18.25	5.37	10.42	15.62	9.58	11.66	11.21	9.39	10.03			

Table 246

	NT2RP1000567	1.21	0.33	0.63	1.21	1.30	2.41	2.12	3.42	1.77		*	+	
5	NT2RP1000574	1.82	0.32	0.03	23.76	28.12	20.34	4.23	4.69	3.79	**	+	**	+
	NT2RP1000577	1.22	0.49	0.73	1.46	1.85	1.75	1.18	2.92	1.35	*	+		
	NT2RP1000579	0.79	0.65	0.57	1.33	1.34	1.32	1.35	2.50	0.76	**	+		
	NT2RP1000581	1.36	0.66	1.82	2.04	1.55	1.78	1.95	2.51	1.03				
	NT2RP1000593	2.64	0.66	1.75	2.65	2.96	1.71	1.41	0.83	1.4				
10	NT2RP1000604	11.50	7.94	7.40	3.94	3.98	3.21	2.12	2.31	2.08	*	-	**	-
	NT2RP1000609	2.53	2.00	0.54	1.02	1.56	1.09	1.82	2.61	1.48				
	NT2RP1000613	1.94	0.88	0.65	1.32	0.99	1.16	0.85	2.58	1.01				
	NT2RP1000622	1.32	0.92	0.99	1.13	1.63	1.80	1.57	3.98	2.19				
	NT2RP1000627	5.47	2.19	3.87	5.94	4.15	4.81	4.23	6.27	4.91				
	NT2RP1000629	1.49	0.86	0.95	1.86	1.84	2.88	2.18	2.88	1.87		*	+	
15	NT2RP1000630	5.89	2.85	5.42	13.99	11.47	13.46	7.36	6.55	7.16	**	+		
	NT2RP1000639	2.68	1.18	0.53	1.84	1.97	0.94	1.56	1.83	1.5				
	NT2RP1000640	81.74	37.60	35.82	57.27	52.32	39.58	48.18	42.34	41.38				
	NT2RP1000646	7.82	4.91	3.97	8.29	9.40	9.31	5.5	5.31	6.52	*	+		
	NT2RP1000659	6.71	2.34	3.90	4.05	6.32	6.12	3.31	4.60	4.15				
20	NT2RP1000674	4.71	2.08	3.93	5.76	7.16	7.25	3.17	4.95	4.5	*	+		
	NT2RP1000677	9.51	6.01	6.41	8.66	8.51	8.83	7.33	7.01	8.68				
	NT2RP1000679	1.23	0.42	0.82	1.73	1.38	1.63	1.09	2.06	0.76	*	+		
	NT2RP1000688	4.67	2.07	2.03	5.85	5.34	3.72	3.1	4.12	2.68				
	NT2RP1000689	2.83	0.64	1.04	1.11	1.67	0.84	1.37	0.88	0.83				
25	NT2RP1000695	1.62	1.12	1.10	1.18	2.39	1.24	1	0.87	0.88				
	NT2RP1000701	0.90	0.82	0.62	0.83	0.27	1.25	0.87	1.19	1.4				
	NT2RP1000702	0.76	0.35	1.53	0.66	1.47	1.82	0.6	1.47	2.57				
	NT2RP1000713	0.23	0.42	0.37	0.34	0.89	0.44	0.17	1.44	1.47				
	NT2RP1000721	10.57	6.36	5.67	7.28	13.00	9.92	8.49	9.05	8.17				
30	NT2RP1000730	2.55	1.65	1.97	4.38	3.90	3.35	1.75	3.65	2.95	*	+		
	NT2RP1000733	4.46	2.99	3.71	5.44	5.04	3.14	1.44	3.93	4.16				
	NT2RP1000738	28.84	10.50	11.79	17.48	18.85	18.44	15.99	11.65	12.72				
	NT2RP1000739	14.40	7.16	8.58	10.60	12.85	8.63	11.15	9.94	11.2				
	NT2RP1000740	3.66	1.37	2.15	2.84	4.09	2.86	2.91	2.60	3.23				
	NT2RP1000746	1.31	0.85	0.82	1.32	1.26	0.89	1.26	2.13	2.46				
35	NT2RP1000750	9.51	4.76	5.09	7.09	6.45	6.48	4.95	5.43	4.72				
	NT2RP1000751	77.49	46.65	53.99	41.34	32.45	28.11	17.67	20.76	21.6		*	-	
	NT2RP1000767	1.53	0.63	1.06	1.68	1.34	1.25	1.21	2.74	2.71				
	NT2RP1000769	4.65	2.64	3.84	2.57	3.18	2.72	4.13	4.77	3.22				
	NT2RP1000780	1.51	0.92	0.80	2.30	1.18	0.64	1.37	0.96	0.77				
40	NT2RP1000782	5.21	2.12	2.72	11.13	10.26	10.71	6.05	7.66	6.54	**	+	*	+
	NT2RP1000796	6.49	4.06	3.11	4.93	5.23	3.73	4.82	3.98	7.13				
	NT2RP1000797	11.72	5.77	5.28	6.51	8.45	5.34	7.81	7.98	9.33				
	NT2RP1000800	0.13	0.54	1.00	1.07	2.16	1.97	0.82	3.18	1.42	*	+		
	NT2RP1000825	3.33	1.37	1.55	2.64	2.23	1.50	1.34	2.32	1.31				
45	NT2RP1000833	6.35	2.68	2.53	4.24	4.98	4.14	2.29	4.43	2.42				
	NT2RP1000834	16.60	5.93	7.79	8.68	7.93	6.33	6.47	7.84	5.03				
	NT2RP1000836	1.43	1.06	0.85	1.19	1.20	0.59	2.19	1.50	0.63				
	NT2RP1000837	6.20	2.33	2.35	4.62	5.53	5.38	4.6	3.52	3.49				
	NT2RP1000846	1.21	0.89	0.89	1.89	2.60	1.73	1.96	1.80	1.08	*	+		
50	NT2RP1000847	2.27	1.79	1.06	1.99	2.12	2.09	2.78	1.80	2.3				
	NT2RP1000851	10.08	6.27	7.87	9.89	12.49	7.13	7.78	9.66	7.43				
	NT2RP1000856	9.90	5.85	7.31	20.58	23.87	20.13	15.75	15.89	19.71	**	+	**	+
	NT2RP1000860	7.91	5.43	8.96	10.11	6.72	7.04	5.54	7.17	4.85				
	NT2RP1000902	2.64	0.85	0.61	5.04	4.02	3.81	3.86	2.25	3.35	*	+		
55	NT2RP1000903	7.75	3.79	2.92	4.96	6.61	5.49	5.15	5.13	5.52				
	NT2RP1000905	3.44	2.09	1.19	3.49	2.21	2.49	3.41	1.72	2.16				
	NT2RP1000915	15.16	7.68	7.64	8.98	6.57	7.27	3.44	4.20	4				

Table 245

	NT2RP1000239	0.54	0.73	0.33	1.02	1.34	0.58	0.16	1.81	1.18			
	NT2RP1000243	0.84	0.90	0.58	2.06	1.73	0.89	0.86	2.41	1.44			
5	NT2RP1000255	0.75	0.34	1.01	1.49	0.71	0.80	0.75	1.85	0.92			
	NT2RP1000259	1.78	1.74	1.10	4.78	3.63	3.84	2.82	4.39	2.57	**	+	+
	NT2RP1000261	1.08	0.77	0.32	2.74	1.80	1.60	0.7	2.51	1.42	*	+	
	NT2RP1000269	12.70	6.05	5.79	12.05	12.78	10.09	7.5	10.31	8.4			
10	NT2RP1000271	65.05	27.46	27.30	118.92	88.05	70.43	44.58	28.04	22.55			
	NT2RP1000272	15.64	8.87	8.62	11.91	10.97	10.04	8.77	5.04	6.08			
	NT2RP1000279	3.64	2.60	2.62	4.01	4.52	4.50	3.4	3.60	2.95	*	+	
	NT2RP1000290	31.80	25.40	25.59	36.52	40.72	40.15	26.39	22.95	29.24	**	+	
	NT2RP1000293	8.90	5.15	6.17	9.07	11.34	10.12	7.62	7.73	8.67			
15	NT2RP1000300	21.75	19.20	18.07	20.53	28.21	20.72	16.45	24.53	12.12			
	NT2RP1000324	12.47	5.32	8.89	10.68	13.57	9.75	6.98	9.83	9.18			
	NT2RP1000325	91.19	35.26	49.60	54.44	61.67	55.26	47.32	30.15	44.99			
	NT2RP1000326	10.60	7.28	6.00	12.46	8.25	10.43	7.71	8.51	5.43			
	NT2RP1000331	13.85	7.24	6.82	12.25	10.31	7.00	5.01	4.72	3.71			
	NT2RP1000333	12.54	6.22	6.09	8.86	8.17	8.74	6.53	7.71	7.88			
20	NT2RP1000336	1.87	1.73	1.02	1.35	1.53	1.21	3.14	2.70	2.83		*	+
	NT2RP1000347	2.75	2.10	2.88	2.09	2.48	2.62	1.53	2.25	0.84			
	NT2RP1000348	1.47	0.48	0.33	1.45	1.42	2.72	1.13	1.89	0.66			
	NT2RP1000349	0.93	0.52	0.64	1.41	1.77	1.72	0.95	0.90	1.19	**	+	
	NT2RP1000353	40.50	18.12	20.02	27.21	16.43	19.17	10.71	8.40	12.57			
25	NT2RP1000356	39.98	22.39	20.90	32.15	26.26	25.06	14.83	10.10	14.28			
	NT2RP1000357	13.61	7.81	6.20	11.20	13.90	12.68	8.98	8.00	11.38			
	NT2RP1000358	11.64	5.39	5.27	10.20	9.77	8.75	7.77	6.88	9.19			
	NT2RP1000360	26.32	15.93	17.17	17.83	19.58	19.99	16.48	15.94	15.67			
	NT2RP1000363	22.05	14.66	16.07	21.39	24.54	24.53	22.26	17.18	17.26			
30	NT2RP1000376	5.84	3.91	5.30	4.51	6.40	6.42	7.18	6.13	5.77			
	NT2RP1000386	31.79	21.04	23.39	64.26	64.31	34.90	56.81	60.95	58.22	*	+	**
	NT2RP1000407	0.29	0.73	0.45	0.62	0.61	0.29	1.08	0.88	0.22			
	NT2RP1000409	2.22	1.91	0.68	2.83	3.38	2.80	2.71	1.86	1.7			
	NT2RP1000413	7.71	3.51	3.63	7.04	7.63	7.01	5.32	4.65	6.75			
35	NT2RP1000416	2.07	0.73	0.71	1.73	2.70	2.64	1.38	1.53	1.42			
	NT2RP1000418	0.88	0.78	0.91	2.07	1.77	2.03	1.84	2.71	1.4	**	+	*
	NT2RP1000420	0.51	0.68	0.34	1.31	0.46	1.21	1.33	1.52	0.65			
	NT2RP1000434	0.66	0.29	2.53	1.80	1.28	1.15	1.63	2.36	0.97			
	NT2RP1000439	13.59	10.41	10.76	8.22	11.99	8.15	6.48	6.20	3.53		*	-
40	NT2RP1000443	1.67	1.60	1.02	3.09	3.95	2.04	3.35	1.76	1.48			
	NT2RP1000447	2.13	0.82	0.90	2.07	1.95	1.21	1.39	1.67	1.12			
	NT2RP1000448	1.39	0.47	0.72	0.68	1.75	1.34	1.82	1.77	0.69			
	NT2RP1000451	5.40	2.45	1.97	5.69	5.15	3.49	1.66	2.36	1.96			
	NT2RP1000458	22.07	12.50	14.79	20.35	29.47	24.03	21.83	19.22	26.03			
	NT2RP1000460	19.74	9.97	12.40	17.61	20.40	21.09	17.72	15.83	18.24			
45	NT2RP1000465	14.77	10.71	12.70	18.32	19.61	21.10	14.71	11.30	11.86	**	+	
	NT2RP1000468	3.47	2.54	4.12	7.07	8.07	7.42	3.93	5.61	4.57	**	+	
	NT2RP1000470	14.45	6.40	6.23	5.28	6.94	7.41	8.62	6.71	6.97			
	NT2RP1000477	0.33	0.76	0.21	0.93	1.49	0.73	0.8	1.04	0.52			
	NT2RP1000478	2.01	1.44	1.12	1.74	1.18	2.18	1.98	3.01	1.97			
50	NT2RP1000481	3.26	1.45	1.19	1.27	1.08	1.24	0.92	2.02	0.85			
	NT2RP1000493	1.13	0.65	0.54	1.16	1.49	1.41	1.57	2.12	0.89	*	+	
	NT2RP1000513	8.57	3.43	5.13	11.73	10.43	8.69	10.51	9.55	9.33			
	NT2RP1000522	9.74	3.47	5.93	6.13	9.61	9.77	8.53	8.00	7.9			
	NT2RP1000533	2.49	0.79	1.93	2.45	2.66	3.02	1.21	2.77	1.5			
	NT2RP1000544	2.42	0.99	0.69	2.39	1.44	1.14	1.43	1.13	2.11			
55	NT2RP1000547	0.17	0.54	0.23	0.77	0.69	0.77	0.43	1.67	0.73	*	+	
	NT2RP1000551	1.62	1.44	0.64	0.50	0.71	0.60	1.24	2.56	1.59			

Table 244

	NT2RM4002535	8.63	5.41	4.92	15.46	13.83	13.63	8.73	8.80	8.02	**	+		
	NT2RM4002554	3.24	2.37	1.91	1.77	3.57	2.77	2.58	2.42	1.39				
5	NT2RM4002558	3.05	3.08	3.12	4.82	3.64	4.78	4.67	4.24	3.5	*	+	*	+
	NT2RM4002565	4.27	2.27	3.74	8.08	6.46	7.43	4.53	4.18	4.36	**	+		
	NT2RM4002567	2.07	1.22	2.13	2.02	9.16	3.14	2.05	3.10	2.71				
	NT2RM4002571	4.37	2.84	3.54	4.69	4.81	3.75	4.16	4.57	3.27				
	NT2RM4002572	6.03	2.28	2.98	6.08	6.68	4.99	9.74	8.44	9.5		*	+	
10	NT2RM4002577	2.75	1.19	0.59	1.71	1.14	1.51	6.59	4.68	6.71		**	+	
	NT2RM4002583	3.95	2.68	2.93	2.91	3.44	3.57	3.57	3.67	3.56				
	NT2RM4002584	6.72	4.52	4.49	7.70	8.13	5.82	4.85	4.09	4.77				
	NT2RM4002593	11.06	6.50	9.20	6.84	5.82	5.78	2.04	3.51	4.47		*	-	
	NT2RM4002594	4.49	2.50	2.60	5.70	6.28	5.59	4.77	7.23	6.06	*	+	*	+
15	NT2RM4002604	4.69	2.15	3.00	3.62	4.47	4.27	3.38	3.51	3.57				
	NT2RM4002614	2.09	1.88	1.83	3.05	2.85	2.71	1.21	3.15	1.87	**	+		
	NT2RM4002616	5.30	2.89	2.15	2.37	1.56	2.52	2.81	1.79	2.9				
	NT2RM4002623	8.57	2.95	4.75	3.25	4.49	3.44	2.87	3.18	2.88				
	NT2RM4002634	1.64	1.74	1.53	1.95	2.12	2.72	2.59	3.50	2.79		**	+	
20	NT2RM4002636	5.12	3.99	4.07	4.89	3.26	2.51	3.1	3.30	2.24		*	-	
	NT2RP1000002	4.91	2.69	3.55	5.37	6.59	6.81	5.02	6.11	5.97	*	+		
	NT2RP1000006	3.58	2.73	3.36	3.30	5.24	3.97	3.46	5.04	3.59				
	NT2RP1000015	0.58	0.54	1.13	1.73	1.75	2.13	1.06	2.60	1.34	**	+		
	NT2RP1000018	0.26	0.38	0.59	1.15	1.19	1.44	1.05	2.21	0.52	**	+		
25	NT2RP1000034	281.35	132.61	141.44	137.16	124.07	106.57	66.03	58.57	54.32				
	NT2RP1000035	3.85	3.38	2.73	3.70	4.44	3.26	2.6	2.77	2.19				
	NT2RP1000040	1.60	1.01	1.16	1.82	1.72	0.90	1.72	1.93	1.4				
	NT2RP1000042	0.16	0.85	0.49	1.42	1.37	0.52	0.89	2.70	1.63				
	NT2RP1000048	3.91	1.94	1.67	2.45	3.78	2.00	3.04	5.80	4.69				
30	NT2RP1000050	2.17	1.06	1.90	2.79	3.16	3.31	1.43	4.06	2.02	*	+		
	NT2RP1000056	29.42	14.22	19.60	15.96	16.06	15.82	8.26	10.94	9.03				
	NT2RP1000058	1.76	1.01	1.59	2.63	1.51	1.74	0.73	1.28	0.3				
	NT2RP1000063	2.86	1.68	1.32	1.33	2.84	1.66	1.17	1.53	1.43				
	NT2RP1000068	2.57	1.65	0.98	2.49	2.52	1.99	1.28	2.09	2.14				
35	NT2RP1000072	111.07	54.80	68.45	57.17	59.96	64.56	51.74	45.59	52.17				
	NT2RP1000073	0.97	0.59	0.56	1.83	1.57	2.36	0.84	2.78	1.72	*	+		
	NT2RP1000078	3.33	1.48	2.67	2.36	2.30	2.50	1.17	3.68	1.39				
	NT2RP1000079	2.67	0.92	1.74	2.69	2.08	2.10	4.5	6.28	4.63		*	+	
	NT2RP1000080	7.28	4.50	5.28	5.11	5.46	5.42	2.3	4.02	3.44				
40	NT2RP1000086	4.35	3.00	3.48	3.24	3.23	2.33	1.02	2.72	1.4		*	-	
	NT2RP1000087	5.00	2.82	2.77	4.73	5.17	3.70	4.25	2.63	3.17				
	NT2RP1000089	21.30	13.02	9.99	15.70	10.56	8.76	7.11	5.03	7.52				
	NT2RP1000090	62.12	34.52	35.37	65.14	57.48	42.93	29.21	27.16	16.48				
	NT2RP1000100	2.17	0.88	1.25	1.24	1.63	1.66	0.75	2.69	2.15				
	NT2RP1000101	6.92	3.86	4.62	6.27	8.56	8.35	6.29	5.31	6.14				
45	NT2RP1000111	3.13	2.02	3.20	4.79	4.46	1.70	2.06	3.98	4.56				
	NT2RP1000112	1.19	1.17	1.40	1.98	2.39	2.90	2.08	3.24	1.09	*	+		
	NT2RP1000124	2.04	1.79	2.18	5.67	6.32	7.61	0.92	3.26	4.08	**	+		
	NT2RP1000125	13.33	6.69	5.55	16.93	13.49	11.53	18.17	14.66	19.62		*	+	
	NT2RP1000129	8.42	3.01	2.92	5.33	4.43	3.32	3.8	3.24	4.62				
50	NT2RP1000130	3.80	3.59	3.16	6.14	5.63	6.01	3.49	3.06	4.37	**	+		
	NT2RP1000154	2.77	1.66	1.73	4.97	6.35	4.78	3.19	4.61	2.92	**	+		
	NT2RP1000163	2.54	1.56	0.69	1.65	3.20	1.85	0.24	3.07	0.88				
	NT2RP1000170	1.25	0.62	0.44	1.93	1.94	1.90	0.89	3.09	1.57	**	+		
	NT2RP1000174	0.77	0.39	0.59	0.80	1.14	0.73	0.83	1.30	0.25				
55	NT2RP1000181	15.66	7.51	13.59	20.37	20.72	18.84	8.95	8.78	5.68	*	+		
	NT2RP1000191	2.05	1.96	1.05	3.54	1.96	2.31	1.34	1.45	2.86				
	NT2RP1000202	1.43	1.24	0.92	2.91	2.20	1.99	0.8	2.37	2.35	*	+		

Table 243

	NT2RM4002189	27.09	13.74	15.74	14.48	15.82	14.29	21.97	17.24	13.45				
	NT2RM4002194	14.06	6.46	6.54	8.20	8.96	5.67	5.24	4.78	7.73				
5	NT2RM4002198	9.72	5.05	4.64	9.60	7.14	7.42	3.99	6.05	4.24				
	NT2RM4002205	6.04	2.24	4.01	10.17	8.07	7.85	3.89	6.45	5.48	*	+		
	NT2RM4002213	8.85	5.39	4.89	8.71	11.13	8.58	6.75	7.49	6.59				
	NT2RM4002216	13.98	11.40	13.83	9.67	12.26	12.25	5.51	6.10	8.64		**	-	
	NT2RM4002226	11.71	3.35	5.45	7.00	6.75	5.32	2.56	2.81	2.06				
10	NT2RM4002237	12.13	5.23	4.66	6.69	6.79	4.62	5.28	4.25	5.13				
	NT2RM4002240	3.83	1.22	1.76	2.57	3.67	3.49	1.94	2.86	3.13				
	NT2RM4002251	4.23	2.41	3.59	5.58	5.63	2.99	3.14	4.22	3.57				
	NT2RM4002256	9.61	4.69	5.30	9.65	8.00	8.72	6.39	6.24	6.69				
	NT2RM4002262	2.51	1.66	3.08	3.94	4.02	2.93	2.54	4.64	4.87				
15	NT2RM4002266	3.81	3.04	1.77	5.13	5.13	3.56	1.74	3.97	3				
	NT2RM4002276	6.07	4.19	4.53	7.03	5.98	6.29	6.42	7.40	4.56				
	NT2RM4002278	5.55	3.50	2.06	5.22	5.68	3.41	2.26	1.92	4.58				
	NT2RM4002281	10.82	3.97	3.78	8.02	12.45	5.87	8.47	7.73	8.03				
	NT2RM4002287	4.73	2.14	2.11	4.48	2.86	2.45	3.19	4.14	1.9				
20	NT2RM4002294	3.56	2.28	1.67	6.99	5.40	3.49	3.08	4.20	3.5				
	NT2RM4002298	4.25	1.83	2.68	5.32	3.86	5.10	6.58	7.80	6.73		**	+	
	NT2RM4002301	2.19	2.10	1.85	3.43	4.22	3.48	1.84	3.94	2.05	**	+		
	NT2RM4002306	4.28	2.89	2.26	4.38	4.65	4.86	3.01	4.34	2.42				
	NT2RM4002323	4.07	3.11	3.95	9.92	6.06	6.87	4.61	4.01	2.32	*	+		
25	NT2RM4002334	48.90	21.85	22.81	35.78	25.59	28.97	30.63	31.70	22.58				
	NT2RM4002339	2.06	1.58	1.46	1.24	1.64	1.38	3.19	1.21	1.93				
	NT2RM4002344	3.34	2.36	2.32	3.06	3.36	3.28	1.98	2.28	1.57				
	NT2RM4002345	3.14	4.48	1.33	2.81	6.18	3.52	3.97	3.81	7.59				
	NT2RM4002352	2.56	1.55	1.37	2.09	1.90	1.83	1.8	2.16	1.75				
30	NT2RM4002362	10.19	5.95	5.50	3.14	3.38	3.88	2.99	2.56	2.32		*	-	
	NT2RM4002373	3.73	2.27	4.81	3.06	4.43	4.48	1.89	2.78	3.21				
	NT2RM4002374	2.46	1.36	2.00	4.92	6.85	2.91	2.01	2.17	2.46				
	NT2RM4002376	3.65	2.05	2.36	5.15	3.88	5.04	5.2	2.97	2.99	*	+		
	NT2RM4002383	5.41	2.46	3.35	8.94	8.52	7.85	5.76	4.08	7.78	**	+		
35	NT2RM4002390	7.22	2.53	2.49	3.89	3.09	3.46	2.47	3.20	2.59				
	NT2RM4002398	4.68	2.42	2.88	5.08	6.85	4.30	3.82	2.28	3.63				
	NT2RM4002409	2.87	2.53	3.04	4.21	5.07	3.80	3.49	3.93	3.64	*	+	*	+
	NT2RM4002414	5.03	1.84	3.97	3.80	4.16	6.28	4.49	4.44	4.68				
	NT2RM4002438	5.21	2.42	2.20	4.07	3.59	4.94	3.44	3.46	2.5				
40	NT2RM4002440	4.95	2.33	3.53	5.69	5.26	3.20	3.34	4.02	4.39				
	NT2RM4002446	6.41	3.72	3.77	5.16	5.23	4.99	5.81	3.91	5.57				
	NT2RM4002450	7.34	5.13	5.19	4.41	3.88	3.16	3.9	3.82	4.13				
	NT2RM4002452	4.76	3.56	2.63	3.31	4.00	4.75	2.58	2.59	2.32				
	NT2RM4002457	3.97	2.35	2.27	5.42	4.08	5.14	4.64	3.85	2.87	*	+		
	NT2RM4002458	2.05	1.17	1.07	1.55	3.27	2.46	2.27	3.06	1.92				
45	NT2RM4002460	1.51	0.73	1.48	0.65	1.16	0.85	1.55	1.39	1.26				
	NT2RM4002464	2.69	1.95	2.48	3.72	3.71	4.31	2.38	2.92	1.83	**	+		
	NT2RM4002479	6.89	5.60	6.27	9.61	8.13	4.62	4.88	6.96	5.42				
	NT2RM4002482	35.61	22.59	16.97	21.71	19.69	20.16	30.11	17.88	24.23				
	NT2RM4002489	15.59	8.96	10.80	10.87	12.64	11.89	10.58	8.12	11.95				
50	NT2RM4002493	3.66	2.45	2.96	3.64	2.32	2.09	3.63	3.17	2.29				
	NT2RM4002499	39.72	27.06	27.17	54.95	67.76	43.13	21.05	20.01	15.47	*	+		
	NT2RM4002504	10.06	5.00	4.83	15.16	13.66	11.30	9.77	10.12	11.17	*	+		
	NT2RM4002506	3.00	2.28	3.10	3.05	3.95	4.66	3.19	3.46	3.27				
	NT2RM4002510	1.71	1.62	1.42	3.05	3.64	3.86	2.57	2.61	2.07	**	+	*	+
55	NT2RM4002527	1.36	1.99	1.93	1.99	2.17	2.01	1.62	2.61	1.13				
	NT2RM4002532	8.36	3.92	4.29	7.17	9.98	8.89	6.69	5.89	6.32				
	NT2RM4002534	5.34	2.37	2.56	3.48	4.24	3.83	3.66	4.16	3.67				

Table 242

	NT2RM4001885	12.23	5.39	5.31	15.89	14.89	14.46	9.96	8.97	11.11	*	+		
	NT2RM4001889	17.90	10.90	9.56	25.74	24.82	26.44	14.72	12.91	12.79	**	+		
5	NT2RM4001894	3.99	3.32	3.07	4.15	4.34	4.16	5.09	3.83	3.49				
	NT2RM4001897	4.68	3.36	3.66	5.57	7.84	6.03	9.17	7.60	6.62	*	+	**	+
	NT2RM4001899	4.37	2.59	2.66	5.10	4.85	5.00	3.8	4.79	3.2	*	+		
	NT2RM4001905	14.13	19.47	18.60	6.62	5.76	7.88	4.18	4.49	4.16	**	-	**	-
	NT2RM4001922	4.57	2.06	2.67	5.98	6.27	5.24	3.2	3.09	2.6	*	+		
10	NT2RM4001930	7.89	5.36	5.01	6.12	7.65	5.79	3.76	3.60	3.88				
	NT2RM4001938	3.35	3.03	2.31	4.03	4.25	3.01	4.12	3.88	3.78		*	+	
	NT2RM4001940	8.88	7.21	7.25	7.65	9.61	6.94	5.41	5.68	5.3		*	-	
	NT2RM4001942	48.53	24.69	36.35	81.10	98.59	62.39	57.57	65.30	79.98	*	+	*	+
	NT2RM4001953	4.86	4.02	3.80	11.16	10.73	8.47	5.44	7.13	6.71	**	+	*	+
15	NT2RM4001965	3.95	3.09	2.78	3.89	4.20	5.02	3.08	4.34	1.87				
	NT2RM4001966	4.92	2.59	2.69	5.18	4.42	3.96	3.32	4.68	3.49				
	NT2RM4001969	4.52	3.56	2.88	4.01	4.54	3.26	3.65	2.05	3.76				
	NT2RM4001974	3.18	2.93	2.68	3.45	3.46	4.29	4	3.93	2.9				
	NT2RM4001979	7.10	5.28	4.65	8.51	9.51	9.19	5.57	5.12	5.65	*	+		
20	NT2RM4001980	8.43	6.53	5.48	9.14	11.80	9.30	5.72	6.09	7.18				
	NT2RM4001984	0.37	0.36	2.68	1.04	2.24	1.27	3.83	2.41	1.54				
	NT2RM4001987	5.43	3.22	4.46	5.44	5.41	4.74	6.11	4.65	5.13				
	NT2RM4002013	4.01	2.99	3.04	5.45	6.17	4.31	4.16	6.39	4.96	*	+		
	NT2RM4002018	1.35	1.30	1.91	4.17	2.80	1.86	2.66	3.82	2.52		*	+	
25	NT2RM4002033	5.95	4.44	3.94	8.70	9.58	8.70	6.99	4.97	5.08	**	+		
	NT2RM4002034	10.16	6.70	5.00	9.69	8.87	7.70	7.22	5.62	6.43				
	NT2RM4002044	17.29	9.91	9.34	16.54	14.23	14.16	9.93	9.20	9.33				
	NT2RM4002047	4.89	3.52	4.39	7.70	9.18	8.38	5.94	5.42	6.2	**	+	*	+
	NT2RM4002054	5.22	3.24	3.62	4.72	4.27	3.95	3.64	4.57	3.02				
30	NT2RM4002055	4.93	3.27	3.62	3.58	4.71	3.15	4.05	4.74	4.4				
	NT2RM4002059	10.05	6.75	9.67	10.16	11.99	13.43	18.25	24.17	33.19		*	+	
	NT2RM4002061	3.42	2.42	3.12	3.99	4.28	3.66	2.26	2.93	1.81	*	+		
	NT2RM4002062	6.37	2.90	3.38	2.10	2.75	3.44	2.98	2.78	3.12				
	NT2RM4002063	8.92	6.28	4.96	9.35	7.20	6.28	7.35	7.35	6.46				
	NT2RM4002066	5.12	2.57	2.72	3.13	3.43	2.84	3.67	3.65	2.97				
35	NT2RM4002067	1.89	1.36	1.11	3.88	3.13	3.49	1.44	3.55	1.91	**	+		
	NT2RM4002073	3.81	3.18	2.17	3.78	3.91	3.14	2.82	4.59	3.46				
	NT2RM4002074	3.75	3.15	4.02	2.89	4.67	3.46	2.89	2.92	2.59		*	-	
	NT2RM4002075	1.30	1.13	1.76	2.76	2.64	2.94	1.69	2.40	1.5	**	+		
	NT2RM4002076	4.00	1.21	3.46	2.32	2.53	2.49	2.84	3.24	1.6				
40	NT2RM4002078	12.66	8.15	5.73	7.75	7.44	9.12	8.77	7.66	8.72				
	NT2RM4002081	5.48	5.00	3.54	7.62	9.31	8.00	5.52	7.35	6.24	**	+		
	NT2RM4002082	4.26	2.31	2.02	3.34	2.38	2.66	2.89	2.98	2.86				
	NT2RM4002093	3.89	2.69	2.12	7.05	6.79	4.47	2.74	4.50	3.5	*	+		
	NT2RM4002109	5.34	3.93	2.60	5.27	7.18	5.20	3.25	3.84	4.24				
45	NT2RM4002115	3.73	2.51	2.56	3.60	4.16	3.32	2.9	3.99	2.74				
	NT2RM4002118	2.39	1.49	2.46	3.46	6.34	3.85	3.47	4.78	5.61		*	+	
	NT2RM4002128	1.76	1.98	1.98	2.53	2.32	2.56	1.95	1.96	1.45	**	+		
	NT2RM4002137	5.40	3.31	3.77	3.32	5.16	4.10	4.08	2.63	2.49				
	NT2RM4002139	6.38	4.93	5.07	14.74	15.06	13.57	6.58	7.18	6.59	**	+		
50	NT2RM4002140	7.07	3.90	5.01	9.78	11.72	9.95	6.8	5.99	6.18	**	+		
	NT2RM4002145	5.69	2.65	3.96	6.30	6.51	4.16	4.2	6.86	5.05				
	NT2RM4002146	12.58	8.18	8.37	8.91	7.31	8.60	4.94	6.93	3.9				
	NT2RM4002161	1.51	1.71	1.05	2.14	2.32	1.65	1.38	2.18	1.6				
	NT2RM4002174	2.04	1.62	2.29	4.40	6.82	5.43	2.41	4.19	3.45	**	+		
	NT2RM4002178	4.27	1.80	4.02	7.72	6.53	7.07	4.59	6.24	4.61	*	+		
55	NT2RM4002180	14.71	6.92	6.30	9.50	9.96	6.78	4.56	4.83	5.69				
	NT2RM4002185	5.31	3.85	4.04	4.39	4.78	3.75	5.7	4.91	5.17				

Table 241

	NT2RM4001597	9.12	5.34	5.09	9.11	10.92	8.47	7.88	8.77	8.89				
	NT2RM4001605	2.56	1.50	0.61	1.85	2.19	2.01	1.99	3.21	1.69				
5	NT2RM4001609	89.25	51.45	54.24	71.13	77.23	52.58	39.95	36.11	41.37				
	NT2RM4001610	12.00	8.23	7.07	12.20	9.44	8.76	11.99	11.53	14.72				
	NT2RM4001611	2.42	1.85	2.60	3.39	3.10	2.31	2.05	3.53	1.91				
	NT2RM4001618	9.99	6.27	7.80	11.85	10.16	11.99	7.45	6.31	7.61				
	NT2RM4001622	26.67	8.64	17.82	10.07	12.08	11.47	11.1	11.92	5.45				
10	NT2RM4001624	6.68	3.27	2.64	4.78	7.08	4.67	4.35	3.33	5.32				
	NT2RM4001625	6.46	4.15	3.63	6.09	6.98	6.57	5.81	6.49	4.68				
	NT2RM4001629	3.08	1.43	1.44	3.13	3.87	3.98	3.34	3.46	2.65				
	NT2RM4001632	29.86	24.78	26.14	43.08	46.42	34.45	16.71	16.75	13.76	*	+	**	-
	NT2RM4001642	2.85	2.24	1.81	3.57	2.70	1.88	1.79	3.45	2.28				
15	NT2RM4001647	17.28	7.78	9.99	11.15	12.30	10.77	8	8.01	6.38				
	NT2RM4001650	0.99	1.51	1.38	2.58	3.80	3.02	1.93	2.32	1.3	**	+		
	NT2RM4001662	7.87	3.75	2.87	5.79	6.00	4.16	5.7	4.40	5.34				
	NT2RM4001666	5.31	2.73	1.99	5.11	5.72	2.91	2.77	3.37	5				
	NT2RM4001670	11.64	5.63	4.93	10.66	7.77	4.83	7.89	5.98	5.85				
20	NT2RM4001682	7.63	4.69	7.88	11.61	13.13	10.67	7.98	7.62	9.49	*	+		
	NT2RM4001710	3.51	1.93	3.14	2.89	2.81	2.52	2.94	3.14	3.23				
	NT2RM4001712	4.09	1.48	2.36	6.28	6.47	3.67	3.14	2.79	2.86				
	NT2RM4001714	9.74	6.27	6.28	8.33	6.94	5.10	4.33	4.54	3.78				
	NT2RM4001715	9.70	6.79	8.58	10.69	5.46	8.50	6.49	7.88	6.36				
25	NT2RM4001727	9.24	3.95	4.64	8.67	8.28	6.42	5.55	4.51	4.54				
	NT2RM4001731	13.05	6.04	4.43	9.34	11.19	3.94	6.46	7.94	7.44				
	NT2RM4001735	10.60	7.33	6.23	6.67	8.99	10.11	4.77	6.71	9.86				
	NT2RM4001739	4.78	4.21	5.14	4.57	4.78	3.04	2.46	4.65	3.94				
	NT2RM4001741	9.97	6.74	4.99	10.67	11.48	8.89	9.93	7.28	7.04				
	NT2RM4001746	4.40	2.92	3.08	6.46	6.23	6.82	4.23	5.87	3.98	**	+		
30	NT2RM4001754	5.88	4.22	4.77	3.77	2.85	3.40	2.26	3.95	2.51	*	-	*	-
	NT2RM4001757	3.98	2.34	2.64	6.30	5.38	5.11	4.27	5.17	3.56	*	+		
	NT2RM4001758	4.03	1.40	1.41	2.95	3.14	0.90	2.11	1.49	2.63				
	NT2RM4001768	9.33	3.18	2.78	8.73	9.23	6.03	4.74	5.46	7.46				
	NT2RM4001775	1.60	0.85	0.48	1.68	1.19	1.13	0.51	1.89	2.16				
35	NT2RM4001776	1.24	0.67	0.70	2.08	1.65	1.01	0.84	1.95	1.26				
	NT2RM4001783	3.30	1.81	1.77	3.52	4.08	2.55	1.62	3.51	1.6				
	NT2RM4001793	5.58	4.64	4.50	8.16	8.15	6.01	4.19	4.76	4.23	*	+		
	NT2RM4001810	3.48	2.21	2.29	3.20	3.69	2.65	2.04	3.39	2.03				
	NT2RM4001813	3.11	0.62	1.16	2.31	2.18	1.56	2	3.91	2.71				
40	NT2RM4001818	3.22	2.40	2.49	5.46	4.70	3.11	4.89	3.44	5.14		*	+	
	NT2RM4001819	11.19	5.78	6.63	9.55	9.42	7.47	10.81	7.51	7.34				
	NT2RM4001823	3.13	1.86	1.29	2.61	3.40	2.11	3.37	1.94	1.66				
	NT2RM4001828	8.26	6.14	6.03	15.07	18.35	11.62	9.17	6.53	11.85	*	+		
	NT2RM4001835	3.34	2.52	2.50	5.07	6.41	5.16	6.93	7.44	8.93	**	+	**	+
45	NT2RM4001836	3.42	2.60	1.50	3.55	5.57	1.89	3.02	2.62	2.83				
	NT2RM4001841	7.03	4.07	5.20	3.69	3.84	5.28	4.46	6.00	6.72				
	NT2RM4001842	2.54	1.03	0.84	4.40	5.14	3.61	2.15	3.10	2.2	*	+		
	NT2RM4001843	7.33	3.08	3.29	4.61	4.36	4.19	6.63	4.29	4.74				
	NT2RM4001856	7.28	3.36	2.92	6.92	6.61	6.34	6.76	4.92	39.96				
	NT2RM4001858	4.41	2.01	2.89	4.19	5.25	3.77	3.99	3.14	3.55				
50	NT2RM4001861	15.16	9.14	7.90	8.10	8.14	9.12	7.69	7.66	6.31				
	NT2RM4001863	5.18	5.03	4.89	5.35	5.57	4.84	4.1	2.95	4.25		*	-	
	NT2RM4001865	4.40	1.50	1.71	4.54	5.77	6.01	3.87	4.69	4.27	*	+		
	NT2RM4001869	6.80	4.12	4.66	5.90	4.78	4.71	3.79	3.46	3.8				
	NT2RM4001873	9.91	7.88	7.75	6.45	7.32	6.28	5.39	4.87	5.18		**	-	
55	NT2RM4001876	20.13	9.94	9.70	9.48	8.26	10.48	13.84	12.07	14.41				
	NT2RM4001890	6.36	4.04	3.70	6.23	5.32	5.66	5.53	4.83	6.28				

Table 240

	NT2RM4001245	8.44	3.64	3.02	4.82	4.98	4.20	6.1	4.52	4.33				
	NT2RM4001247	3.08	1.77	1.70	4.91	4.46	4.61	4.28	3.78	3.75	**	+	*	+
5	NT2RM4001256	2.68	1.97	1.44	2.55	3.37	2.58	2.83	3.46	2.47				
	NT2RM4001258	3.01	1.08	-1.54	2.58	2.80	3.08	2.91	3.65	2.07				
	NT2RM4001267	3.85	1.81	3.09	2.74	2.85	2.20	1.95	2.54	1.48				
	NT2RM4001273	4.22	3.00	2.18	5.27	4.13	4.07	4.07	4.58	3.5				
	NT2RM4001281	4.83	2.17	2.72	3.21	3.18	3.70	4.92	3.04	3.76				
10	NT2RM4001286	200.90	135.14	135.42	284.75	209.56	246.97	164.2	134.29	151.3	*	+		
	NT2RM4001290	9.86	4.80	5.69	5.57	5.18	5.39	8.08	8.05	9.32				
	NT2RM4001309	4.86	3.06	2.25	4.98	6.28	4.18	3.55	4.91	3.92				
	NT2RM4001313	5.02	3.13	3.38	10.23	11.21	8.30	5.64	5.09	6.07	**	+		
	NT2RM4001316	3.10	1.87	1.63	4.90	3.32	2.72	2.34	3.07	2.48				
15	NT2RM4001320	3.57	1.99	1.80	4.35	3.95	2.99	2.67	3.38	1.95				
	NT2RM4001321	2.36	1.76	2.19	4.88	3.23	3.63	2.96	3.26	2.18	*	+		
	NT2RM4001325	4.26	2.86	2.43	3.61	4.06	3.37	3.66	2.87	3.97				
	NT2RM4001333	9.63	4.30	7.26	19.73	18.36	12.94	10.99	11.48	14.86	*	+		
	NT2RM4001340	15.08	7.81	6.58	8.67	7.96	8.93	6.1	7.09	9.47				
20	NT2RM4001344	5.69	1.98	2.69	4.58	3.47	5.21	3.57	4.25	3.42				
	NT2RM4001347	2.27	2.16	1.78	2.66	5.15	3.34	3.4	3.43	2.43		*	+	
	NT2RM4001357	6.92	4.15	5.35	6.32	6.10	5.55	4.34	5.12	6.64				
	NT2RM4001360	5.77	3.29	3.38	4.26	4.44	4.12	4.69	3.72	3.64				
	NT2RM4001371	4.54	2.79	3.83	7.15	6.45	5.83	3.62	4.03	2.04	*	+		
25	NT2RM4001377	10.12	5.47	3.83	5.72	6.90	5.90	6.53	6.36	7.54				
	NT2RM4001382	27.64	18.16	15.30	26.18	25.29	24.42	17.41	14.13	18.42				
	NT2RM4001384	2.18	1.75	1.21	2.08	4.07	2.57	1.73	1.84	2.63				
	NT2RM4001400	1.97	1.68	1.05	5.11	4.43	3.04	4.16	3.64	2.67	*	+	*	+
	NT2RM4001409	2.47	2.29	2.32	4.11	6.40	4.45	3.11	3.39	3.96	*	+	*	+
30	NT2RM4001410	3.95	1.97	3.57	4.82	7.04	5.31	4.02	3.43	5.37	*	+		
	NT2RM4001411	0.83	0.77	0.89	2.84	2.80	2.65	2.26	2.50	1.14	**	+		
	NT2RM4001412	3.72	2.65	2.59	3.20	3.12	4.78	3.05	4.81	2.12				
	NT2RM4001414	4.96	2.76	1.91	3.88	3.24	3.95	8.58	4.11	4.61				
	NT2RM4001436	10.71	5.74	4.93	8.68	8.18	5.45	5.99	5.69	6.31				
	NT2RM4001437	3.31	2.10	1.69	4.84	3.86	4.25	2.81	3.12	5.09	*	+		
35	NT2RM4001444	17.08	11.93	9.02	14.24	23.31	17.28	9.91	10.57	15.43				
	NT2RM4001454	1.52	1.27	0.92	2.66	2.43	3.18	2.53	2.75	3.91	**	+	*	+
	NT2RM4001455	1.97	1.35	0.94	1.41	2.43	2.26	1.92	2.49	2.53				
	NT2RM4001483	8.15	6.48	6.83	17.59	20.73	16.59	7.89	9.09	9.1	**	+		
	NT2RM4001489	2.71	2.11	2.58	4.94	4.32	3.30	3.82	3.97	2.42	*	+		
40	NT2RM4001495	18.14	8.14	7.60	6.61	8.27	8.97	13.02	9.27	7.52				
	NT2RM4001499	12.77	8.16	6.92	3.39	3.00	2.48	3.08	3.42	2.67	*	-	*	-
	NT2RM4001515	3.27	1.91	1.68	2.35	4.06	1.83	1.52	2.44	1.37				
	NT2RM4001519	5.12	2.84	4.04	2.41	3.33	2.32	2.38	4.57	1.4				
	NT2RM4001522	6.04	4.16	3.86	10.17	8.78	6.98	5.57	5.11	4.64	*	+		
45	NT2RM4001523	2.87	2.23	1.80	2.40	4.75	2.55	2.53	3.39	1.48				
	NT2RM4001550	9.31	4.21	5.82	7.65	10.18	9.65	4.79	5.78	4.65				
	NT2RM4001553	13.10	6.91	9.72	15.17	15.42	12.48	9.84	10.30	8.03				
	NT2RM4001554	6.26	1.91	2.23	3.10	3.46	2.19	2.03	3.40	3.47				
	NT2RM4001557	1.82	1.50	1.72	2.44	4.16	3.37	2.15	2.77	2.22	*	+	*	+
50	NT2RM4001565	4.45	2.55	3.09	4.16	3.19	4.16	3.34	4.45	3.44				
	NT2RM4001566	8.15	6.36	5.54	21.07	22.32	19.38	14.82	13.59	12.47	**	+	**	+
	NT2RM4001569	1.07	2.72	1.12	1.58	1.44	1.53	1.39	2.06	0.92				
	NT2RM4001579	2.12	1.63	1.82	2.74	2.69	2.72	4.53	3.33	2.15	**	+		
	NT2RM4001582	2.62	2.33	2.55	3.71	4.48	4.20	3.06	3.87	3.26	**	+	*	+
55	NT2RM4001589	8.35	5.09	6.66	12.13	12.37	10.11	11.51	12.65	14.42	*	+	**	+
	NT2RM4001592	3.41	2.19	1.04	2.79	0.97	1.51	1.07	1.30	2.99				
	NT2RM4001594	6.13	3.39	4.24	4.38	6.50	3.46	3.95	3.79	5.84				

Table 239

	NT2RM4000841	5.06	3.39	2.43	4.05	5.93	4.27	4.63	4.44	4.32				
5	NT2RM4000846	9.09	5.94	7.28	12.84	12.70	15.11	9.96	10.78	11.36	**	+	*	+
	NT2RM4000848	7.88	5.40	5.25	6.98	11.06	5.33	7.53	6.82	8.25				
	NT2RM4000852	6.75	4.64	5.34	13.69	17.70	14.08	11.97	13.34	10.93	**	+	**	+
	NT2RM4000855	4.73	2.86	4.28	6.84	5.05	6.75	4.95	4.71	3.45	*	+		
	NT2RM4000859	13.33	7.63	8.66	12.33	11.71	13.85	10.92	13.05	13.48				
10	NT2RM4000868	3.39	2.48	3.24	2.56	3.27	2.72	2.54	2.52	2.34				
	NT2RM4000870	7.43	4.59	4.58	4.56	7.18	4.83	5.21	5.55	10.16				
	NT2RM4000879	5.36	4.71	2.54	2.94	5.60	3.69	4.73	3.05	8.38				
	NT2RM4000882	13.28	7.67	8.34	13.87	16.02	12.84	11.37	9.53	8.64				
	NT2RM4000887	7.73	5.89	6.66	6.98	5.77	6.42	10.56	10.15	7.39				
	NT2RM4000895	5.73	3.47	4.08	7.64	7.37	6.94	4.46	6.14	5.95	*	+		
15	NT2RM4000897	7.53	4.28	4.64	9.70	11.04	6.20	7.51	8.32	7.28				
	NT2RM4000901	2.04	1.85	1.79	2.60	2.63	3.31	2.13	2.92	1.47	*	+		
	NT2RM4000950	0.56	0.78	1.17	2.14	1.27	1.24	1.41	2.19	1.17				
	NT2RM4000965	9.86	4.20	4.55	3.73	5.50	4.12	5.03	3.46	4.87				
	NT2RM4000971	5.30	5.00	2.48	7.54	6.04	2.89	3.53	4.64	7.17				
20	NT2RM4000979	4.99	2.53	1.69	2.02	3.14	2.85	2.38	2.83	3.27				
	NT2RM4000987	2.44	1.53	2.68	3.20	3.68	2.75	2.62	4.83	3.64				
	NT2RM4000989	4.94	3.38	3.37	4.04	2.94	2.51	3.27	4.13	3.58				
	NT2RM4000991	0.93	1.02	1.31	2.15	2.31	2.55	2.33	4.87	2.11	**	+		
	NT2RM4000992	11.24	7.63	10.16	7.25	5.43	5.90	4.54	4.54	4.18	*	-	**	-
25	NT2RM4000996	4.06	2.34	3.75	9.54	9.91	8.12	3.46	4.48	3.87	**	+		
	NT2RM4000997	9.49	3.35	2.92	6.90	7.64	7.96	5.25	6.12	5.29				
	NT2RM4001001	22.10	15.26	10.21	12.02	9.69	11.49	22.6	17.92	9.97				
	NT2RM4001002	5.24	3.19	3.25	8.21	8.99	8.70	5.14	6.05	8.69	**	+		
	NT2RM4001016	4.56	3.14	3.04	3.93	5.46	2.92	3.16	3.93	3.9				
30	NT2RM4001025	115.98	53.32	70.45	58.33	60.27	42.54	40.15	40.87	41.74				
	NT2RM4001027	0.14	0.43	0.68	0.22	0.31	0.94	0.68	1.67	1.36				
	NT2RM4001032	1.80	1.46	0.81	3.10	2.87	2.32	1.9	2.71	1.77	*	+		
	NT2RM4001047	1.37	0.95	0.95	2.05	2.61	2.62	1.72	2.11	1.51	**	+	*	+
	NT2RM4001049	10.71	3.63	3.82	6.40	6.54	4.49	5.52	5.09	5.26				
35	NT2RM4001051	6.70	3.93	4.20	7.11	12.15	4.54	5.61	4.11	11.9				
	NT2RM4001052	8.14	4.27	4.08	6.07	7.39	5.45	8.57	7.89	6.02				
	NT2RM4001053	27.19	14.20	21.35	17.33	19.31	15.07	12.02	9.63	10.5				
	NT2RM4001054	3.61	1.72	2.96	2.73	3.57	4.09	2.66	3.55	3.62				
	NT2RM4001059	7.61	4.52	5.00	8.40	9.15	6.24	6.45	6.67	8.15				
40	NT2RM4001071	4.06	2.69	2.57	4.40	6.02	4.14	3.25	5.00	2.66				
	NT2RM4001084	4.94	2.76	3.04	3.73	6.30	5.46	4.17	4.56	4.31				
	NT2RM4001092	7.29	2.48	2.72	5.06	4.22	4.55	3.22	2.32	2.04				
	NT2RM4001100	12.18	6.64	7.67	10.87	11.09	10.86	6.95	8.94	8.4				
	NT2RM4001116	1.86	1.58	1.69	2.27	2.62	2.03	2.58	1.98	1.6	*	+		
	NT2RM4001119	4.12	2.84	2.77	3.79	5.02	3.34	2.23	3.61	4.07				
45	NT2RM4001140	16.77	10.70	11.39	11.80	11.74	11.76	7	6.89	6.74		*	-	
	NT2RM4001148	13.85	6.50	6.41	8.02	8.87	5.20	9.72	12.16	8.38				
	NT2RM4001151	3.04	2.82	2.68	3.38	3.91	4.17	3.34	5.07	4.04	*	+		
	NT2RM4001155	3.85	1.95	2.51	2.48	2.96	3.06	2.88	3.51	1.43				
	NT2RM4001157	4.58	2.01	1.48	3.42	3.84	2.43	3.68	3.71	2.97				
50	NT2RM4001160	6.16	2.57	2.15	5.06	4.60	3.14	2.68	2.65	4.39				
	NT2RM4001163	28.46	18.93	15.30	35.95	37.53	27.65	20.27	18.39	15.85				
	NT2RM4001187	5.15	3.42	2.71	6.56	6.27	4.41	3.87	4.84	4.54				
	NT2RM4001191	4.08	1.58	2.81	4.80	3.69	3.67	1.67	2.71	2.13				
	NT2RM4001200	5.87	3.23	4.14	11.90	10.51	10.62	3.7	6.92	6.13	**	+		
	NT2RM4001203	5.49	3.54	4.23	5.75	6.16	5.89	3.38	6.37	4.68				
55	NT2RM4001204	1.21	0.66	1.10	1.28	1.38	1.01	0.49	2.21	0.81				
	NT2RM4001217	2.79	2.05	1.40	2.03	2.19	2.27	3.13	2.64	2.28				

Table 238

	NT2RM4000514	5.53	2.38	2.75	8.23	11.94	5.81	3.95	5.11	3.73			
	NT2RM4000515	16.72	6.51	7.89	17.68	19.19	15.60	8.65	8.97	10.58			
5	NT2RM4000517	52.07	29.36	32.93	47.60	48.78	40.92	19.63	19.22	17.95			
	NT2RM4000520	2.37	1.45	1.44	1.17	1.70	1.58	0.83	2.01	2.17			
	NT2RM4000531	1.99	2.27	1.67	2.66	3.68	3.90	3.09	4.12	3.6*	+	**	+
	NT2RM4000532	1.32	0.65	0.82	1.96	2.81	1.58	1.14	2.83	2.21*	+		
	NT2RM4000533	3.05	2.29	3.20	1.70	2.71	1.77	1.32	2.54	1.44			
10	NT2RM4000534	1.94	0.89	1.21	1.63	2.79	1.54	1.47	2.29	1.5			
	NT2RM4000563	8.72	3.55	3.49	6.44	4.79	3.51	5.01	4.09	5.24			
	NT2RM4000566	4.57	2.22	2.28	4.38	4.92	2.84	2.28	2.65	3.1			
	NT2RM4000568	3.97	2.58	1.85	3.65	4.45	3.11	2.68	3.32	5.31			
	NT2RM4000585	4.60	2.16	1.71	2.71	3.64	3.29	2.11	2.49	3.12			
15	NT2RM4000587	2.44	1.07	2.17	2.90	3.56	2.74	2.55	3.03	3.48			
	NT2RM4000590	2.10	1.53	1.91	1.79	3.35	2.47	1.66	3.17	1.73			
	NT2RM4000593	7.87	4.39	5.71	12.59	12.23	10.68	4.27	7.40	6.4**	+		
	NT2RM4000595	2.17	1.55	2.08	3.28	3.82	2.26	2.16	3.13	4.82			
	NT2RM4000683	12.55	6.66	4.52	9.64	5.73	7.27	4.44	2.74	3.84			
20	NT2RM4000611	4.28	4.40	1.85	3.49	3.25	3.20	4.67	2.74	3.19			
	NT2RM4000616	3.34	2.92	1.37	4.32	4.33	3.69	3.56	2.86	2.97			
	NT2RM4000621	16.48	12.72	9.94	21.48	22.15	19.49	9.4	8.56	8.57*	+		
	NT2RM4000648	2.01	1.43	1.11	2.32	1.95	1.99	1.76	2.62	1.65			
	NT2RM4000649	5.47	3.71	4.22	6.21	6.35	6.84	6.07	5.86	5.42*	+		
25	NT2RM4000658	8.60	4.07	5.16	8.70	7.92	4.74	5.84	5.98	5.36			
	NT2RM4000661	10.99	4.92	5.69	11.11	10.38	8.21	15.64	14.68	17.57	*	+	
	NT2RM4000673	9.96	5.23	4.31	6.63	5.66	5.28	8.2	4.95	5.83			
	NT2RM4000674	5.01	2.88	2.93	4.58	4.03	4.02	5.28	3.25	4.19			
	NT2RM4000689	6.44	3.20	3.50	4.50	6.19	4.47	3.52	4.05	3.79			
30	NT2RM4000698	35.87	22.93	21.16	15.46	17.90	22.28	17.5	16.82	14.8			
	NT2RM4000700	3.46	2.08	2.83	3.85	2.02	2.52	2.49	2.37	1.32			
	NT2RM4000701	9.78	5.90	5.74	10.46	14.71	8.86	7.95	6.35	8.32			
	NT2RM4000712	2.69	1.64	2.42	4.68	4.33	3.64	2.57	3.33	2.41*	+		
	NT2RM4000717	12.02	5.07	6.36	11.87	8.62	8.11	7.27	6.28	7.15			
35	NT2RM4000733	8.98	3.57	6.27	6.72	6.26	7.78	7.76	4.90	6			
	NT2RM4000734	9.72	3.11	3.90	7.75	4.13	5.58	5.8	4.00	5.07			
	NT2RM4000741	4.49	2.29	3.56	3.14	3.42	3.32	3.44	4.03	2.18			
	NT2RM4000744	3.69	2.68	2.61	2.80	6.32	4.46	2.85	3.92	3			
	NT2RM4000749	11.40	7.45	11.83	11.62	13.08	12.36	13.08	12.48	13.4			
40	NT2RM4000751	6.54	4.81	4.52	15.28	14.53	10.59	6.43	6.81	9.13**	+		
	NT2RM4000752	4.53	2.37	3.48	4.41	5.68	4.78	3.23	4.75	8.68			
	NT2RM4000760	4.53	2.84	2.99	5.14	6.37	2.91	5.41	3.73	5.34			
	NT2RM4000761	996.52	787.70	799.46	925.45	928.88	688.98	521.4	1076.26	1043			
	NT2RM4000764	27.63	19.80	15.48	20.84	20.29	16.92	30.21	26.08	33.56			
45	NT2RM4000768	14.67	8.26	9.77	8.91	9.00	6.52	3.2	6.21	5.06	*	-	
	NT2RM4000778	4.92	2.41	4.01	2.84	3.65	2.97	1.85	2.67	2.07			
	NT2RM4000779	8.60	6.98	9.29	9.01	13.32	14.40	9.71	7.63	14.65			
	NT2RM4000787	4.24	2.50	3.69	7.64	7.50	6.95	5.13	4.57	3.51**	+		
	NT2RM4000790	3.29	2.32	3.49	4.70	4.95	5.71	2.8	3.89	2.61*	+		
	NT2RM4000795	17.99	8.62	8.95	7.60	7.29	5.12	10.59	11.21	13.05			
50	NT2RM4000796	9.52	5.97	4.89	6.98	7.91	6.65	7.34	5.94	6.5			
	NT2RM4000798	4.86	3.32	1.92	4.08	3.21	6.07	3.4	3.16	3.56			
	NT2RM4000800	25.53	16.14	15.27	24.04	32.78	23.66	18.49	15.32	20.57			
	NT2RM4000813	9.68	4.14	5.79	3.70	5.76	3.65	5.28	6.71	6.8			
	NT2RM4000820	6.65	4.53	5.35	8.29	7.69	8.43	5.66	5.99	4.55*	+		
55	NT2RM4000827	7.32	3.89	5.09	8.78	8.63	9.18	6.29	5.93	5.97*	+		
	NT2RM4000830	6.10	3.43	4.84	5.47	6.76	7.08	5.66	6.54	4.35			
	NT2RM4000833	7.52	4.61	4.22	4.98	5.08	4.81	5.7	5.33	4.23			

Table 237

	NT2RM4000155	8.41	4.25	5.85	5.71	7.89	3.63	6.31	2.89	10.88			
	NT2RM4000156	4.06	2.82	3.12	3.91	5.14	3.25	4.15	3.34	7.54			
5	NT2RM4000167	2.76	1.86	2.44	3.27	3.78	2.46	1.7	2.54	2.08			
	NT2RM4000169	19.79	11.82	12.59	15.78	28.83	16.15	10.62	8.62	22.74			
	NT2RM4000191	5.46	2.93	3.98	7.00	12.87	3.95	5.75	4.22	5			
	NT2RM4000197	6.21	3.61	5.57	1.78	3.32	3.20	2.07	2.72	3.62			
	NT2RM4000198	6.32	5.24	5.02	9.16	10.86	8.33	6.38	7.74	6.83	**	+	
10	NT2RM4000199	3.97	1.83	1.79	3.99	4.05	3.81	2.77	3.05	3.55			
	NT2RM4000200	3.35	2.42	1.54	4.45	2.14	1.95	1.94	2.20	2.16			
	NT2RM4000202	3.63	1.09	1.43	2.56	2.87	2.44	2.2	2.07	1.78			
	NT2RM4000210	4.14	2.52	2.72	3.86	8.22	3.80	3.01	2.97	3.68			
	NT2RM4000215	5.18	3.07	5.47	7.27	8.45	5.15	4.83	4.89	4.29			
15	NT2RM4000220	2.94	2.54	2.79	4.64	4.57	3.49	4.1	4.60	5.77	*	+	*
	NT2RM4000229	5.01	3.09	3.00	5.45	4.41	4.69	4.07	4.71	4.56			
	NT2RM4000231	4.55	4.22	5.24	5.48	9.85	6.48	5.25	5.36	6.29			
	NT2RM4000233	15.69	9.94	12.92	10.36	8.30	6.63	11.95	12.79	13.03			
	NT2RM4000244	3.55	2.12	1.68	2.06	1.74	1.35	2.28	2.40	1.4			
20	NT2RM4000251	3.33	1.28	1.28	2.48	6.47	3.24	2.39	3.65	3.7			
	NT2RM4000255	2.86	2.35	2.55	3.65	4.00	4.45	3.46	3.56	4.05	**	+	**
	NT2RM4000265	4.79	2.78	4.25	9.35	12.26	8.62	3.89	4.46	7.5	**	+	
	NT2RM4000283	70.67	47.66	58.69	22.90	27.64	23.33	20.04	20.53	29.33	**	-	**
	NT2RM4000284	3.79	2.43	3.13	4.73	5.37	4.18	3.75	4.06	5.01	*	+	
25	NT2RM4000290	3.63	2.15	2.31	4.25	6.01	4.45	4.22	4.40	5.11	*	+	*
	NT2RM4000295	2.18	1.74	1.84	1.64	1.85	1.54	2.16	2.51	2.05			
	NT2RM4000306	9.76	5.69	5.53	3.29	5.79	3.80	4.99	4.91	4.19			
	NT2RM4000307	1.99	1.95	1.34	6.27	6.75	5.25	9.66	12.35	13.1	**	+	**
	NT2RM4000309	4.39	2.45	3.20	3.45	3.57	3.25	2.21	2.77	3.12			
30	NT2RM4000313	4.53	2.93	3.37	6.76	7.38	6.57	4.37	4.56	4.95	**	+	
	NT2RM4000318	3.24	1.42	3.10	6.35	5.08	6.14	3.2	4.49	3.95	**	+	
	NT2RM4000324	3.33	2.91	2.72	5.10	4.10	4.09	3.41	4.13	3.13	*	+	
	NT2RM4000326	2.66	2.08	2.02	2.52	2.48	2.90	1.91	4.16	2.37			
	NT2RM4000327	5.98	3.83	5.87	11.13	9.36	9.04	5.82	4.08	6.84	**	+	
35	NT2RM4000344	18.32	6.89	6.35	13.95	16.21	14.72	10.48	11.38	12.84			
	NT2RM4000349	6.58	3.84	3.66	6.40	5.99	6.38	4.94	4.61	4.8			
	NT2RM4000354	5.00	2.70	3.37	3.28	2.86	2.19	2.4	2.57	3.45			
	NT2RM4000356	4.16	1.61	1.73	2.39	4.18	5.03	2.81	3.86	2.82			
	NT2RM4000366	51.05	23.81	40.37	61.56	72.80	50.45	36.85	39.74	37.86			
	NT2RM4000368	4.89	2.95	4.56	12.45	6.89	8.75	3.93	5.00	5.04	*	+	
40	NT2RM4000373	3.91	2.54	3.44	5.84	6.63	5.55	3.15	4.00	4.07	**	+	
	NT2RM4000386	2.58	1.67	2.32	2.56	2.07	2.16	1.54	2.11	1.77			
	NT2RM4000395	7.43	3.02	3.38	5.38	8.33	4.62	3.75	3.62	2.41			
	NT2RM4000414	8.01	4.62	4.45	4.72	6.23	4.47	6.27	6.74	7.44			
	NT2RM4000417	3.81	2.15	2.35	2.45	4.44	3.37	1.96	2.94	5.43			
45	NT2RM4000421	4.32	3.14	3.21	5.52	5.35	4.92	3.28	3.80	2.75	*	+	
	NT2RM4000425	5.83	3.82	4.77	12.15	12.72	12.65	6.8	7.68	8.72	**	+	*
	NT2RM4000433	3.24	1.87	2.39	3.27	3.60	3.54	5.12	3.76	4.32		*	+
	NT2RM4000436	5.20	2.98	5.09	4.80	5.70	3.50	3.27	3.38	2.58			
	NT2RM4000444	2.77	3.48	2.67	4.83	3.05	2.47	3.64	2.66	2.6			
50	NT2RM4000457	15.74	6.60	7.42	19.58	21.46	13.99	7.15	6.46	8.49			
	NT2RM4000471	2.61	2.36	2.45	4.93	5.40	4.25	2.75	2.88	3.1	**	+	*
	NT2RM4000472	18.08	9.02	12.03	39.03	47.24	22.44	10.92	7.97	22.46	*	+	
	NT2RM4000486	3.65	3.27	2.92	5.46	7.04	5.93	3.74	4.42	3.79	**	+	
	NT2RM4000490	4.88	4.19	2.87	2.63	4.80	4.02	3.72	6.17	4.86			
55	NT2RM4000496	4.08	3.13	4.22	2.96	3.44	3.46	3.56	4.42	3.51			
	NT2RM4000505	13.63	8.59	11.63	16.76	17.74	14.63	12.93	15.48	13.33	*	+	
	NT2RM4000511	58.96	34.63	49.12	52.44	54.53	48.10	20.11	21.96	24.23		*	-

Table 236

	NT2RM2002492	21.46	13.29	16.96	23.80	28.37	23.64	14.79	12.77	15.74	*	+		
5	NT2RM2002575	14.83	8.83	9.60	12.39	15.50	9.64	5.47	4.77	4.26		*	-	
	NT2RM2002580	10.54	5.71	6.88	9.66	15.19	13.67	6.89	8.73	8.24				
	NT2RM2002592	21.59	13.02	21.47	22.05	25.36	18.29	13.81	13.13	16.44				
	NT2RM2002608	14.51	10.47	15.10	11.85	17.10	10.74	12.25	12.95	16.2				
	NT2RM2002615	7.16	4.68	6.11	4.32	3.11	3.23	2.34	3.30	2.9	*	-	*	-
10	NT2RM2002622	7.42	4.82	9.06	37.13	40.07	28.33	10.87	12.05	11.06	**	+	*	+
	NT2RM2002630	7.98	5.03	5.96	13.25	13.42	12.82	6.17	6.79	6.95	**	+		
	NT2RM2002634	5.03	2.59	3.78	7.49	9.33	4.95	4.93	3.29	2.99				
	NT2RM2002645	23.59	12.83	21.14	22.24	21.50	17.33	18.84	24.20	13.44				
	NT2RM2002646	14.00	9.34	10.97	13.76	16.22	12.07	10.73	9.69	15.13				
	NT2RM2002647	20.09	9.61	14.48	15.78	21.02	13.76	11.26	13.21	13.26				
15	NT2RM2002652	5.04	3.66	3.21	6.10	6.51	3.39	2.65	3.93	4.06				
	NT2RM2002692	7.77	5.58	7.47	11.71	20.77	13.98	11.29	13.53	12.54	*	+	**	+
	NT2RM2002721	24.72	15.21	18.70	28.40	46.33	25.49	20.79	20.54	32.86				
	NT2RM2002748	79.54	53.04	79.10	79.94	75.77	78.90	26.04	31.33	37.42		*	-	
	NT2RM2002764	5.43	3.03	2.52	10.76	7.77	4.58	3.86	3.77	3.77				
20	NT2RM2002772	11.93	7.88	8.81	11.61	12.84	7.73	4.61	5.99	7.99				
	NT2RM2002811	9.63	5.90	5.86	8.67	8.08	6.76	5.99	5.21	6.14				
	NT2RM2002818	6.94	3.95	3.88	7.36	7.54	5.33	2.65	4.06	4.31				
	NT2RM2002879	2.57	1.77	2.32	2.29	3.75	1.84	3.18	4.11	4.37		*	+	
	NT2RM2002979	11.80	7.84	8.67	10.47	13.00	9.87	8.38	6.63	6.92				
25	NT2RM2002981	4.75	2.96	3.25	4.20	5.55	4.27	4.3	5.20	4.19				
	NT2RM2002995	3.40	2.64	2.64	3.84	3.50	4.10	2.62	3.34	2.85	*	+		
	NT2RM2003031	3.92	1.02	1.63	4.33	4.68	2.72	3.7	2.74	3.72				
	NT2RM2003042	21.41	10.74	8.21	17.59	19.62	15.87	7.89	8.90	9.64				
	NT2RM2003044	3.74	2.06	1.81	3.99	6.41	3.64	2.33	3.97	3.12				
30	NT2RM2003090	4.60	2.18	1.89	2.49	4.89	3.16	3.07	3.31	2.92				
	NT2RM2003095	3.67	1.54	1.20	3.30	4.47	3.32	3.18	3.65	3.25				
	NT2RM2003116	5.36	5.13	6.83	5.86	7.80	6.25	3.24	6.72	6.31				
	NT2RM2003222	2.53	2.08	1.54	2.39	2.31	1.74	0.73	3.10	1.35				
	NT2RM2003224	15.53	10.87	13.94	24.44	25.63	15.64	6.09	8.22	11.35				
35	NT2RM2003250	14.48	5.65	5.15	9.14	10.21	4.29	3.99	3.24	3.21				
	NT2RM2003258	2.29	2.33	1.33	2.70	2.97	1.92	4.64	2.60	3.37				
	NT2RM2003262	12.60	10.45	8.76	10.06	13.00	11.50	9.36	7.15	7.82				
	NT2RM4000023	1.99	1.44	1.54	4.90	4.52	3.88	4.13	2.29	4.66	**	+		
	NT2RM4000024	2.91	2.48	1.20	3.30	4.50	2.17	2.67	1.90	2.29				
40	NT2RM4000027	8.53	4.07	5.06	2.82	3.04	1.62	1.79	2.08	2.61				
	NT2RM4000030	5.84	5.94	5.16	8.87	6.03	4.15	5.42	5.51	5.41				
	NT2RM4000033	1.51	1.27	1.03	2.93	3.16	1.42	1.59	1.08	1.27				
	NT2RM4000034	2.39	1.22	1.22	3.53	2.94	1.45	2.28	1.04	1.5				
	NT2RM4000046	2.68	1.77	1.53	3.42	3.11	1.75	3.04	1.82	3.01				
	NT2RM4000052	4.15	1.71	1.72	3.48	3.49	2.40	3.28	1.71	3.37				
45	NT2RM4000054	26.80	19.29	17.31	21.55	22.04	23.11	25.09	20.51	27.5				
	NT2RM4000061	2.10	1.10	0.99	1.68	1.71	1.22	2.51	1.77	1.98				
	NT2RM4000074	9.55	7.34	6.67	13.37	15.17	7.83	6.11	5.94	7.27				
	NT2RM4000085	2.96	0.88	2.51	4.65	4.96	3.33	2.05	2.94	4.07				
	NT2RM4000086	5.73	3.89	4.54	5.27	5.35	3.12	1.65	3.66	5.45				
50	NT2RM4000100	5.36	2.82	2.66	5.25	5.01	3.76	5.82	4.53	4.32				
	NT2RM4000101	3.85	2.50	2.70	3.14	2.31	2.97	4.25	3.04	4.67				
	NT2RM4000102	36.64	21.10	21.71	40.33	40.26	40.80	33.11	25.16	36.34				
	NT2RM4000104	1.41	0.89	0.77	2.16	1.98	1.38	2.39	2.42	1.99		**	+	
	NT2RM4000115	1.25	1.28	1.23	1.59	2.32	1.33	1.87	1.24	1.33				
	NT2RM4000129	2.55	2.06	1.92	3.48	3.51	3.33	3.04	3.19	2.54	**	+		
55	NT2RM4000139	2.48	1.32	1.75	2.52	1.95	2.96	1.63	2.77	2.58				
	NT2RM4000149	1.92	1.98	1.88	3.18	3.67	2.07	1.43	1.95	2.57				

Table 235

	NT2RM2001887	4.05	2.53	2.07	3.94	3.93	2.72	2.74	2.00	2.87			
	NT2RM2001896	968.51	557.14	625.69	446.49	419.99	290.65	817.5	613.90	955.7			
5	NT2RM2001902	1.32	1.09	1.03	2.63	3.33	2.08	2.5	1.84	1.37*	+		
	NT2RM2001903	10.52	8.17	6.65	10.52	9.78	8.75	7	6.78	10.05			
	NT2RM2001930	5.61	3.44	3.21	5.48	6.96	3.46	4.44	4.85	6			
	NT2RM2001935	3.82	1.91	1.54	3.50	4.79	3.97	2.7	3.75	4.62			
	NT2RM2001936	5.82	4.45	4.35	6.11	7.15	5.56	4.64	4.90	5.38			
10	NT2RM2001939	8.71	5.44	6.44	8.93	8.81	3.78	2.77	3.30	4.35	*	-	
	NT2RM2001941	6.75	2.80	2.92	6.78	5.32	3.44	5.9	3.69	5.46			
	NT2RM2001950	7.11	3.51	4.45	5.50	5.26	4.20	5.45	4.64	5.47			
	NT2RM2001952	2.47	1.60	2.55	2.69	4.21	2.27	1.88	1.01	2.57			
	NT2RM2001976	28.42	15.82	19.71	28.96	35.93	24.29	16.42	13.99	23.68			
15	NT2RM2001982	4.42	1.68	2.40	3.83	3.46	2.37	2.4	2.21	2.73			
	NT2RM2001983	2.90	2.45	2.37	3.29	3.84	2.68	3.58	3.72	3.62	**	+	
	NT2RM2001984	9.80	5.19	8.10	8.76	9.27	5.57	9.18	6.75	8.16			
	NT2RM2001989	11.11	6.20	6.87	11.27	9.42	7.93	6.29	5.35	7.09			
	NT2RM2001996	14.80	9.47	8.75	13.23	9.98	7.81	6.58	6.93	7.66			
20	NT2RM2001997	6.28	4.07	2.81	7.04	8.03	5.28	7.41	5.47	7.79			
	NT2RM2001998	4.75	3.45	3.00	4.75	6.36	4.13	5.37	3.71	5.85			
	NT2RM2001999	10.41	5.56	7.08	6.38	11.36	7.48	5.73	5.79	10.27			
	NT2RM2002003	10.66	5.49	8.27	9.09	11.29	8.39	10.04	6.40	24.73			
	NT2RM2002004	1.63	1.64	2.11	1.09	1.63	1.85	1.23	1.86	1.25			
25	NT2RM2002009	4.47	4.69	3.31	8.66	11.16	6.73	5.88	6.79	8.4*	+	*	+
	NT2RM2002014	2.01	1.63	2.37	3.01	3.07	2.13	1.7	1.98	2.36			
	NT2RM2002019	24.72	12.04	19.38	13.08	13.17	13.22	11.49	8.63	11.15			
	NT2RM2002029	6.40	7.22	6.06	8.84	11.57	6.10	8.68	6.47	10.53			
	NT2RM2002030	5.25	5.14	4.68	5.36	8.72	3.88	5.86	5.43	6.29			
30	NT2RM2002034	8.15	6.62	4.89	14.77	20.00	13.04	13.54	8.03	15.03*	+		
	NT2RM2002049	3.95	2.79	2.89	4.72	8.26	6.22	5.53	3.64	6.92*	+		
	NT2RM2002055	0.27	0.82	0.37	0.80	1.13	1.85	1.04	1.68	0.63			
	NT2RM2002072	15.43	11.44	16.71	17.13	17.10	21.32	19.05	15.56	22.41			
	NT2RM2002088	7.49	4.56	5.69	7.90	6.52	5.70	5.75	6.67	7.06			
35	NT2RM2002091	15.11	10.25	9.22	22.42	19.93	19.66	8.6	12.53	10.62*	+		
	NT2RM2002100	4.63	3.56	2.83	7.24	10.07	3.66	3.27	4.23	5.16			
	NT2RM2002109	5.17	3.65	3.18	8.12	10.78	4.99	4.99	4.26	6.51			
	NT2RM2002126	17.67	11.99	12.06	15.99	24.43	15.73	17.49	13.92	19.27			
	NT2RM2002128	3.48	2.83	1.99	3.84	5.46	3.66	3.24	2.92	3.02			
	NT2RM2002129	4.13	2.91	3.80	6.20	6.87	4.06	5.78	4.67	7.21			
40	NT2RM2002142	9.10	5.41	12.04	10.00	15.48	9.23	8.42	6.45	11.18			
	NT2RM2002144	3.36	3.30	2.97	3.37	3.35	3.00	3.79	3.97	3.53	*	+	
	NT2RM2002145	6.78	4.33	5.19	6.26	8.85	5.46	5.35	5.34	6.65			
	NT2RM2002153	23.74	16.73	21.12	12.42	16.25	18.91	6.63	5.66	6.16	**	-	
	NT2RM2002163	3.16	2.77	2.30	3.73	2.93	2.52	3.34	5.43	2.91			
45	NT2RM2002170	3.33	3.09	3.14	5.55	7.02	5.69	2.89	3.62	2.63**	+		
	NT2RM2002178	5.79	2.91	3.21	5.77	6.57	4.62	3.87	4.53	5.9			
	NT2RM2002179	2.75	2.13	3.45	13.46	15.53	10.86	9.37	9.17	14.68**	+	*	+
	NT2RM2002270	6.01	3.32	3.61	5.54	5.51	3.68	4.91	5.60	3.82			
	NT2RM2002326	3.03	1.98	3.43	11.14	9.52	7.64	7.73	8.34	6.08**	+	**	+
50	NT2RM2002337	4.10	3.34	2.03	4.41	8.58	3.20	3.05	2.79	3.57			
	NT2RM2002339	7.43	4.86	4.58	4.19	4.70	6.27	6.54	7.31	8.68			
	NT2RM2002345	4.47	3.51	3.00	6.85	4.79	5.38	4.35	5.79	4.74			
	NT2RM2002368	4.40	3.36	3.81	8.23	7.04	7.08	3.82	5.20	3.26**	+		
	NT2RM2002381	1.63	1.57	2.71	2.99	3.95	2.46	1.73	3.26	3.15			
55	NT2RM2002424	6.30	4.83	5.88	15.99	15.30	14.85	6.59	9.16	9.11**	+		
	NT2RM2002450	4.28	2.58	3.43	4.26	4.94	3.98	2.13	3.11	1.67			
	NT2RM2002482	3.24	2.34	3.46	4.41	2.79	3.35	3.25	3.29	2.2			

Table 234

	NT2RM2001664	5.67	1.74	2.53	4.44	5.25	4.88	2.57	2.77	3.63				
	NT2RM2001668	7.83	4.11	5.80	12.91	11.03	10.32	6.9	5.64	6.89	*	+		
5	NT2RM2001670	5.07	2.93	3.57	4.21	4.81	3.14	3.67	3.52	4.76				
	NT2RM2001671	2.26	2.13	2.75	6.03	4.05	5.74	4.08	5.29	5.9	*	+	**	+
	NT2RM2001675	0.53	0.71	0.81	1.96	1.15	1.66	0.84	1.96	0.76	*	+		
	NT2RM2001681	1.11	1.22	1.01	3.34	4.29	2.27	1.69	3.16	1.72	*	+		
	NT2RM2001685	3.03	2.26	1.29	2.06	2.47	1.90	1.92	3.02	2.65				
10	NT2RM2001688	2.78	1.66	2.54	4.45	4.23	2.30	3.29	2.37	2.72				
	NT2RM2001695	7.30	3.32	3.64	20.95	20.35	18.16	12.07	10.51	12.36	**	+	**	+
	NT2RM2001696	13.28	6.12	3.86	8.81	10.82	9.78	6.65	6.44	6.65				
	NT2RM2001698	8.16	4.37	3.88	5.88	6.34	6.37	6.66	7.84	5.32				
	NT2RM2001699	2.40	2.32	1.42	3.33	3.59	3.21	1.64	3.47	3.24	*	+		
15	NT2RM2001700	2.41	1.38	1.03	2.93	2.03	1.36	1.5	2.70	2.35				
	NT2RM2001704	6.94	4.34	5.63	17.99	22.84	16.16	12.13	13.06	13.82	**	+	**	+
	NT2RM2001706	5.19	2.60	4.07	6.98	8.64	6.26	3.29	5.55	5.04	*	+		
	NT2RM2001714	1.72	1.75	2.15	3.05	3.64	2.23	1.71	2.86	4.84				
	NT2RM2001716	16.89	6.66	8.99	10.52	14.37	8.33	10.03	10.20	4.62				
20	NT2RM2001718	13.66	7.01	6.41	14.04	11.83	11.25	5.12	7.15	10.53				
	NT2RM2001723	6.13	3.06	3.78	9.65	9.89	7.73	4.12	5.35	3.42	*	+		
	NT2RM2001727	5.93	4.01	4.52	4.87	5.62	5.99	5.45	6.14	6.96				
	NT2RM2001730	3.02	1.57	1.66	3.08	3.68	4.44	2.79	3.79	2.6				
	NT2RM2001738	6.78	3.40	5.60	4.93	5.41	3.52	4.55	4.50	4.55				
25	NT2RM2001743	4.12	2.65	1.97	3.64	5.10	2.62	3.21	3.25	2.82				
	NT2RM2001753	4.87	2.89	3.87	7.06	7.68	7.46	4.96	5.77	5.73	**	+		
	NT2RM2001755	11.15	5.43	7.63	8.83	12.50	9.88	7.94	7.72	5.34				
	NT2RM2001760	6.52	3.36	4.22	8.42	9.37	6.40	10.28	9.84	11.76			**	+
	NT2RM2001765	2.13	1.98	1.79	3.23	3.97	3.48	2.65	2.41	2.82	**	+	*	+
30	NT2RM2001767	12.87	8.82	9.72	11.08	15.03	8.12	9.19	9.22	14.64				
	NT2RM2001768	3.41	2.58	3.68	3.47	6.28	4.04	2.49	2.74	3.01				
	NT2RM2001771	4.11	3.62	4.50	11.05	14.86	9.39	5.06	5.82	8.71	**	+		
	NT2RM2001778	1.70	1.61	1.19	3.14	4.69	2.67	2.01	2.74	1.97	*	+		
	NT2RM2001782	3.37	2.78	3.39	3.01	4.99	4.13	3.83	4.97	5.07		*	+	
35	NT2RM2001784	3.64	1.97	1.45	2.55	4.38	1.85	2.15	2.16	2.26				
	NT2RM2001785	11.40	5.25	4.67	8.49	7.03	6.72	4.99	4.72	4.92				
	NT2RM2001792	5.79	3.39	4.17	6.69	5.40	4.24	3.59	5.22	5.39				
	NT2RM2001795	9.85	4.56	3.32	7.91	9.48	5.77	7.27	6.25	5.93				
	NT2RM2001797	5.04	2.64	2.13	7.82	15.93	10.34	3.54	4.95	3.54	*	+		
40	NT2RM2001800	3.26	2.51	2.46	4.20	4.38	3.21	2.99	3.72	2.42				
	NT2RM2001803	3.60	2.31	2.65	4.14	6.89	5.00	2.04	3.10	3.17				
	NT2RM2001805	1.03	0.92	2.17	2.21	3.99	1.67	0.87	3.16	1.79				
	NT2RM2001806	5.77	1.94	1.66	4.46	3.73	2.85	3.42	3.44	3.44				
	NT2RM2001813	3.38	1.75	1.74	2.55	3.99	2.42	1.83	1.59	3.71				
45	NT2RM2001814	3.09	1.71	2.83	3.06	4.28	2.96	1.96	3.02	3.47				
	NT2RM2001818	2.38	1.33	1.54	3.40	2.50	2.32	1.89	3.32	1.89				
	NT2RM2001823	1.26	1.12	0.39	0.95	1.88	0.91	0.96	2.06	1.08				
	NT2RM2001825	10.44	6.78	7.32	10.86	11.22	8.43	10.54	16.17	16.27		*	+	
	NT2RM2001832	4.52	2.18	1.93	4.11	5.31	3.71	1.98	4.92	3.68				
	NT2RM2001839	16.50	9.01	12.64	20.38	24.01	12.42	26.45	40.89	49.99		*	+	
50	NT2RM2001840	7.75	3.07	2.83	17.33	13.18	10.65	7.84	6.84	7.97	*	+		
	NT2RM2001851	7.34	4.30	5.43	12.87	13.61	10.39	7.97	4.60	6.49	**	+		
	NT2RM2001855	5.55	3.48	2.68	4.96	5.56	4.20	7.12	5.33	6.29				
	NT2RM2001867	3.35	3.93	2.06	5.68	4.49	3.19	3.59	2.29	4.09				
	NT2RM2001869	28.84	23.52	26.51	34.13	35.79	25.24	15.38	12.75	18.43			**	-
55	NT2RM2001879	0.65	1.04	0.48	2.26	1.68	1.53	1.35	1.12	1.72	*	+	*	+
	NT2RM2001883	3.25	3.47	2.90	9.89	19.82	12.52	5.43	4.02	7.24	*	+		
	NT2RM2001886	2.86	1.25	2.11	3.84	5.06	2.09	1.79	1.91	1.93				

Table 233

	NT2RM2001201	13.08	8.55	9.63	10.72	11.46	9.31	8.37	10.02	9.79			
	NT2RM2001221	6.92	2.79	3.15	5.91	7.22	4.72	4.61	5.13	3.98			
5	NT2RM2001238	2.81	1.05	1.43	3.40	2.72	2.10	1.81	2.65	3.37			
	NT2RM2001243	6.98	4.99	5.16	9.29	9.00	6.32	4.34	5.08	4.64			
	NT2RM2001244	4.98	5.59	4.41	14.49	19.11	7.34	5.11	6.41	7.87			
	NT2RM2001247	15.41	9.79	11.87	12.82	15.98	10.20	6.66	8.32	9.67			
	NT2RM2001256	2.93	2.70	3.12	2.39	2.54	2.02	2.24	3.49	2.22	*	-	
10	NT2RM2001269	1.76	1.73	1.47	3.07	6.49	3.10	1.39	5.05	2.29			
	NT2RM2001278	7.64	6.14	6.38	12.27	11.97	10.88	6.39	7.92	7.27	**	+	
	NT2RM2001291	4.14	2.35	1.90	4.62	4.03	2.79	3.65	2.48	3.16			
	NT2RM2001294	10.67	6.20	5.16	12.58	9.68	9.06	8.36	5.49	6.33			
	NT2RM2001295	4.70	3.78	3.23	5.43	4.66	4.21	4.46	4.14	4.92			
15	NT2RM2001302	5.63	4.69	4.19	1.74	2.61	0.97	2.97	3.64	4.24	**	-	
	NT2RM2001306	2.52	1.56	1.39	3.47	5.32	4.74	2.64	2.44	2.72	*	+	
	NT2RM2001312	1.22	1.12	0.35	2.84	2.71	1.41	1.03	2.09	1.77			
	NT2RM2001319	5.09	3.21	4.08	5.71	5.46	5.01	3.84	5.43	5.66			
	NT2RM2001324	8.85	3.42	3.83	7.05	8.29	8.06	5.36	6.31	4.89			
20	NT2RM2001345	12.36	6.03	4.96	4.58	10.06	7.26	10.14	5.50	8.05			
	NT2RM2001360	9.69	4.48	4.35	8.36	5.80	5.82	6.45	4.63	6.16			
	NT2RM2001370	1.53	1.04	0.81	1.70	1.86	1.18	1.6	2.44	2.47		*	+
	NT2RM2001391	1.02	1.38	1.05	3.81	3.30	1.71	1.72	1.73	1.75	*	+	**
	NT2RM2001393	6.61	4.78	7.01	5.53	6.68	4.32	4.86	4.39	4.92			
25	NT2RM2001420	2.35	0.95	1.41	3.00	4.15	1.59	1.98	2.45	1.71			
	NT2RM2001423	11.93	5.27	6.94	5.59	7.80	3.34	2.15	4.14	4.71			
	NT2RM2001424	18.20	9.15	9.42	11.35	10.96	8.30	11.11	9.35	12.67			
	NT2RM2001482	15.21	7.55	7.78	14.57	12.13	9.92	11.31	8.31	11.15			
	NT2RM2001499	16.92	9.02	7.05	8.26	6.45	6.32	5.19	4.43	7.42			
30	NT2RM2001504	3.91	2.51	1.97	4.23	4.34	3.86	4.03	2.84	4.42			
	NT2RM2001524	2.28	1.47	1.87	2.95	3.08	2.80	2.63	3.34	2.29	*	+	
	NT2RM2001530	0.78	0.43	0.54	2.16	2.44	1.43	1.65	1.93	1.93	*	+	**
	NT2RM2001533	5.77	3.13	3.08	6.59	7.98	5.62	5.57	5.84	5.16			
	NT2RM2001540	29.91	19.29	20.03	25.11	24.66	12.51	8.93	9.56	11.82		*	-
	NT2RM2001544	5.22	2.70	2.16	5.77	5.72	5.39	4.13	3.93	3.57			
35	NT2RM2001547	10.18	3.47	3.29	5.82	9.93	4.61	8.42	7.52	11.22			
	NT2RM2001558	4.96	2.25	2.36	3.07	3.85	4.04	4.67	2.71	4.49			
	NT2RM2001575	4.76	2.31	3.04	7.85	7.43	4.47	3.66	3.23	5.49			
	NT2RM2001582	3.25	3.39	2.40	5.42	5.69	4.66	5.53	3.88	4.63	**	+	*
	NT2RM2001588	2.97	1.41	1.47	4.20	4.38	3.50	3.05	3.37	3.85	*	+	
40	NT2RM2001592	1.95	2.06	1.67	3.66	3.58	2.66	2.98	2.38	2.72	*	+	*
	NT2RM2001603	7.68	4.12	5.42	8.07	9.92	5.79	4.3	6.45	7.62			
	NT2RM2001605	6.36	3.57	2.87	8.10	9.32	7.63	6.11	4.82	7.04	*	+	
	NT2RM2001611	4.43	2.58	2.01	5.92	8.58	4.85	5.15	3.69	4.23			
	NT2RM2001613	5.87	2.94	3.70	6.48	9.87	7.29	8.01	9.10	11.64		*	+
45	NT2RM2001626	11.27	5.06	6.34	7.63	9.90	5.32	10.52	8.42	10.76			
	NT2RM2001632	8.60	4.62	8.41	12.48	14.32	11.43	11.18	11.01	12.96	*	+	*
	NT2RM2001633	1.62	1.36	1.29	4.23	3.15	3.36	2.92	3.34	2.97	**	+	**
	NT2RM2001635	6.76	5.69	4.97	6.78	9.41	8.26	7.11	8.37	8.13		*	+
	NT2RM2001636	4.43	3.06	3.83	4.52	5.10	3.16	3.81	3.42	3.49			
50	NT2RM2001637	2.79	1.78	2.31	4.20	4.67	2.96	3.03	3.93	2.28	*	+	
	NT2RM2001639	4.58	2.65	2.19	3.05	3.54	3.49	2.88	2.16	3.34			
	NT2RM2001641	3.30	2.69	1.81	2.72	3.01	1.92	1.93	2.95	2.91			
	NT2RM2001643	3.00	1.41	2.34	4.92	3.73	2.89	3.15	3.35	3.43			
	NT2RM2001648	3.60	1.94	2.50	5.96	6.92	4.37	6.52	6.11	8.21	*	+	**
	NT2RM2001652	4.13	2.45	1.80	4.68	5.72	3.86	2.23	2.96	4.29			
55	NT2RM2001659	1.81	1.41	1.26	2.31	1.88	1.34	2.02	2.75	2.32		*	+
	NT2RM2001660	2.12	1.41	1.99	2.87	4.60	2.23	2.13	3.03	3.35			

Table 232

	NT2RM2000658	7.80	7.19	11.39	10.12	9.62	7.80	6.87	5.57	7.23				
	NT2RM2000660	27.64	11.87	13.50	20.31	25.06	18.91	13.54	13.65	15.55				
5	NT2RM2000669	7.79	4.71	4.17	9.97	13.43	8.55	3.67	4.50	6.66				
	NT2RM2000689	29.82	30.60	28.82	42.51	72.34	55.67	22.11	19.71	38.62	*	+		
	NT2RM2000691	4.67	3.54	3.74	5.23	6.41	4.14	4.29	3.98	4.19				
	NT2RM2000714	13.27	8.60	10.19	9.82	10.81	9.42	13.37	9.65	17.53				
	NT2RM2000718	1.36	1.54	1.09	3.28	7.10	3.02	2.42	2.48	2.19		**	+	
10	NT2RM2000732	6.10	4.20	5.69	12.72	15.74	11.49	5.7	6.42	7.79	**	+		
	NT2RM2000735	24.38	15.21	20.46	56.19	49.62	47.05	16.37	24.66	27.14	**	+		
	NT2RM2000740	6.48	2.95	2.62	6.53	5.49	3.44	3.93	3.46	2.74				
	NT2RM2000743	21.35	12.67	14.35	10.73	9.73	9.68	2.24	1.81	2.16		**	-	
	NT2RM2000772	11.89	7.81	9.52	17.15	14.77	14.45	6.23	7.95	10.31	*	+		
15	NT2RM2000773	11.75	6.40	6.69	9.73	11.32	9.29	9.82	8.51	8.01				
	NT2RM2000776	12.66	6.48	11.36	17.08	19.56	14.42	12.22	8.19	11.56	*	+		
	NT2RM2000784	11.22	7.09	6.83	7.88	10.63	6.42	6.22	6.90	7.64				
	NT2RM2000795	9.52	5.29	6.34	17.74	18.61	15.80	6.53	8.43	10.09	**	+		
	NT2RM2000796	27.57	17.52	26.46	2.02	2.40	3.17	1.82	2.65	1.66	**	-	**	-
20	NT2RM2000798	14.84	8.16	10.91	45.29	27.47	24.14	26.69	20.97	28.82	*	+	**	+
	NT2RM2000801	37.70	23.20	28.38	26.35	37.85	28.51	31.37	32.22	38.5				
	NT2RM2000821	3.67	2.04	2.27	8.85	6.90	6.15	5.86	5.63	5.4	**	+	**	+
	NT2RM2000829	36.66	22.85	41.47	29.93	25.94	16.17	15.48	17.92	19.23		*	-	
	NT2RM2000837	5.77	3.15	3.99	6.12	6.76	5.46	5.15	4.55	4.39				
25	NT2RM2000924	6.69	5.13	4.70	12.18	14.72	8.21	5.5	6.80	8.89	*	+		
	NT2RM2000930	14.27	7.36	9.58	15.72	15.41	13.15	7.93	7.73	11.49				
	NT2RM2000937	2.93	2.09	3.52	5.00	4.64	3.14	1.89	3.58	2.8				
	NT2RM2000939	6.56	3.88	4.32	5.94	7.25	6.23	4.34	5.73	5.56				
	NT2RM2000942	141.00	79.29	113.17	107.50	122.19	108.41	73.07	66.91	67.18				
30	NT2RM2000951	4.09	2.69	2.78	3.88	3.40	4.39	3.48	3.83	3.33				
	NT2RM2000952	5.14	3.58	3.50	6.02	4.82	4.48	3.55	3.67	3.9				
	NT2RM2000966	11.75	10.12	10.87	9.00	11.41	11.06	9.18	10.30	5.82				
	NT2RM2000973	22.49	16.16	17.58	24.24	28.57	21.97	14.32	17.17	15.94				
	NT2RM2000983	10.51	6.87	10.06	15.15	16.05	11.81	9.62	13.40	12.47	*	+		
	NT2RM2000984	3.34	2.49	1.94	4.17	6.33	3.91	3.14	3.09	3.89				
35	NT2RM2000994	17.72	5.91	15.58	25.00	22.32	16.64	8.13	8.32	6.15				
	NT2RM2001004	6.95	4.49	3.43	6.09	8.10	5.86	5.16	4.92	6.83				
	NT2RM2001022	113.50	66.21	87.63	148.44	181.02	157.90	78.72	73.28	91.6	*	+		
	NT2RM2001035	10.78	6.86	10.47	14.95	15.69	13.90	7.29	8.73	9.42	*	+		
	NT2RM2001038	4.09	2.22	2.89	6.55	5.43	6.97	3.62	3.51	3.32	*	+		
40	NT2RM2001043	2.10	1.71	2.70	4.88	5.53	4.13	3.52	4.59	4.54	**	+	*	+
	NT2RM2001050	8.66	4.61	6.50	7.54	9.45	9.85	5.61	5.16	6.52				
	NT2RM2001055	4.62	4.14	3.41	6.16	5.15	5.46	4.13	4.67	4.8	*	+		
	NT2RM2001065	6.07	2.63	3.08	8.01	7.85	5.22	3.46	3.40	2.98				
	NT2RM2001075	101.53	60.27	56.87	59.75	60.87	48.63	40.45	36.79	33.7				
45	NT2RM2001083	13.68	8.75	8.30	8.28	9.63	7.55	10.14	7.92	7.18				
	NT2RM2001100	8.62	6.13	5.38	7.77	11.80	7.92	8.12	5.53	6.12				
	NT2RM2001105	18.36	12.31	11.09	26.95	28.47	25.34	13.8	13.76	12.91	**	+		
	NT2RM2001109	5.91	3.28	4.91	5.36	6.02	5.10	4.49	5.78	4.86				
	NT2RM2001110	9.13	5.14	5.81	7.93	9.23	8.95	5.5	5.40	7.45				
50	NT2RM2001126	4.23	4.04	4.69	10.78	10.09	7.28	4.72	4.73	5.55	**	+		
	NT2RM2001131	9.35	4.34	5.26	6.74	7.12	6.44	4.76	4.21	3.9				
	NT2RM2001141	9.27	7.43	7.38	17.22	17.01	12.50	8.53	7.69	8.62	*	+		
	NT2RM2001152	3.64	1.47	1.46	2.09	3.47	2.45	1.42	1.89	2.81				
	NT2RM2001177	8.38	4.92	5.00	10.70	11.58	8.68	5.53	6.90	6.04	*	+		
55	NT2RM2001194	10.76	6.38	8.60	11.33	15.08	9.90	8.38	9.20	9.42				
	NT2RM2001195	3.62	3.00	3.18	4.85	7.13	3.67	3.45	3.91	3.64				
	NT2RM2001196	7.18	4.57	6.50	10.22	17.76	12.85	4.98	6.12	9.71	*	+		

Table 231

	NT2RM2000250	6.85	2.87	3.45	6.74	6.95	8.42	4.64	4.72	5.57				
	NT2RM2000259	9.6	4.08	4.77	6.02	9.47	7.13	5.19	6.42	6.9				
5	NT2RM2000260	9.93	9.2	6.51	4.88	7.9	8.73	11.23	8.04	9.57				
	NT2RM2000265	2.4	1.14	0.66	1.28	0.86	1.86	1.3	1.27	1.08				
	NT2RM2000287	10.73	4.68	6.12	10.38	10.35	12.59	6.93	10.27	8.06				
	NT2RM2000306	16.48	15.91	13.02	16.75	16.33	10.75	17.88	8.38	16.11				
	NT2RM2000312	57.19	46.28	42.21	59.66	41.08	60.14	43.74	21.02	32.47				
10	NT2RM2000322	6.45	2.73	3.3	5.49	4.98	2.77	3.63	4.55	3.78				
	NT2RM2000343	5.35	4.3	5.69	10.01	10.47	9.81	6.04	4.74	6.91	**	+		
	NT2RM2000359	5.94	2.73	3.95	5.3	4.77	4.66	3.08	3.35	2.77				
	NT2RM2000362	15.37	16.06	11.14	15.03	19.07	17.41	12.3	11.08	9.04				
	NT2RM2000363	2.27	1.12	1.53	3.15	1.57	1.39	1.27	1.95	1.06				
15	NT2RM2000368	20.14	10.44	9.67	11.84	14.77	11.87	10.3	9.5	10.03				
	NT2RM2000371	111	74.6	73.79	116.3	62.15	121.3	50.3	42.75	56.01		*	-	
	NT2RM2000374	4.78	2.52	1.94	6.65	5.32	5.42	4.66	3.93	3.68	*	+		
	NT2RM2000387	11.91	6.37	5.79	20.24	13.27	20.63	9.51	12.58	11.14	*	+		
	NT2RM2000393	3.45	1.01	1.83	2.71	1.61	3.18	1.81	3.53	1.53				
20	NT2RM2000395	1.44	0.49	0.91	2.24	0.76	1.26	1.08	2.52	0.72				
	NT2RM2000402	7.26	1.87	2.95	6.33	6.77	7.71	5.51	6.64	5.38				
	NT2RM2000405	5.34	2.42	2.76	3.26	3.78	4.88	2.25	2.56	2.19				
	NT2RM2000407	19.34	9.57	10.6	5.59	9.51	9.38	8.65	7.51	10.04				
	NT2RM2000410	3.06	1.14	0.97	2.09	2.96	2.28	2.57	1.94	2.16				
25	NT2RM2000420	4.52	1.56	1.71	6.72	7.81	5.85	4.96	3.72	3.6	*	+		
	NT2RM2000422	14.32	4.96	7.79	15.68	12.45	9.99	14.38	10.45	10.29				
	NT2RM2000423	3.93	2.29	3.18	9.3	10.31	11.58	4.01	3.67	2.37	**	+		
	NT2RM2000452	4.1	1.67	3.69	10.71	9.43	9.18	6.96	4.45	5.45	**	+		
	NT2RM2000469	1.22	0.59	0.27	2.22	1.54	1.32	1.52	1.06	1.82				
30	NT2RM2000490	4.98	2.59	1.93	4.39	4.04	3.10	5.95	3.52	4.92				
	NT2RM2000497	2.77	1.77	1.58	7.44	5.74	5.87	2.86	3.26	4.3	**	+		
	NT2RM2000502	4.18	2.99	2.68	7.32	4.36	3.54	3.69	2.68	5.35				
	NT2RM2000504	2.49	1.56	2.01	5.06	3.93	4.92	5.83	4.60	4.88	**	+	**	+
	NT2RM2000514	5.60	3.19	3.45	8.34	7.66	5.47	4.66	4.70	6.69				
35	NT2RM2000522	0.63	0.58	0.61	1.36	0.80	1.01	0.53	0.67	1.87				
	NT2RM2000540	5.03	4.07	2.80	5.25	6.86	2.78	4.31	3.32	4.3				
	NT2RM2000556	0.38	0.75	0.50	1.40	1.96	0.69	3.19	0.77	0.73				
	NT2RM2000565	4.89	2.53	3.37	4.40	4.50	4.25	5.66	3.06	4.57				
	NT2RM2000566	5.85	4.38	3.46	8.37	5.27	4.67	4.65	4.38	5.92				
	NT2RM2000567	4.29	3.05	2.89	4.78	3.00	1.68	3.19	2.38	4.64				
40	NT2RM2000569	6.50	3.15	2.85	8.65	8.54	6.48	4.57	3.91	4.43				
	NT2RM2000577	11.83	4.68	6.45	6.50	8.99	3.96	4.84	6.67	8.79				
	NT2RM2000581	6.47	3.33	5.21	7.46	8.40	4.99	4.74	5.34	7.76				
	NT2RM2000582	5.88	3.81	3.49	9.44	7.98	6.09	7.69	6.61	8.15	*	+	*	+
	NT2RM2000588	22.92	13.30	11.99	23.97	16.17	19.54	16.8	11.46	18.28				
45	NT2RM2000589	11.18	6.26	6.74	9.54	8.57	7.04	5.39	6.22	7.18				
	NT2RM2000594	11.31	9.59	11.31	3.91	4.21	3.25	3.27	3.48	2.37	**	-	**	-
	NT2RM2000599	22.01	15.12	17.66	19.78	24.09	21.66	15.2	13.93	15.24				
	NT2RM2000609	2.49	1.70	2.43	4.47	3.94	3.24	1.96	3.24	2.3	*	+		
	NT2RM2000612	3.82	2.55	2.84	4.46	7.55	3.95	4.78	3.73	4.27				
50	NT2RM2000622	8.85	7.06	10.37	13.55	16.80	10.42	7.83	8.92	10.48				
	NT2RM2000623	23.78	13.14	15.60	19.91	22.53	23.26	22.22	15.23	19.45				
	NT2RM2000624	16.48	10.64	4.76	11.37	17.12	10.30	8.76	8.23	137.7				
	NT2RM2000632	5.44	2.83	2.35	3.85	3.76	2.79	2.22	2.21	7.42				
	NT2RM2000635	2.91	2.32	2.35	7.82	9.57	5.76	5	4.36	4.83	**	+	**	+
	NT2RM2000636	3.87	2.82	3.19	5.69	5.77	3.68	4.63	3.86	4.55				
55	NT2RM2000639	4.56	3.86	3.29	4.47	7.45	4.02	3.6	6.93	4.67				
	NT2RM2000649	4.09	4.81	3.74	4.86	8.90	5.64	4.74	5.82	6.57				

Table 230

NT2RM1000956	16.88	9.05	9.11	8.8	11.37	15.79	15.38	17.86	10.86				
NT2RM1000960	13.57	6.62	8.78	22.97	30.24	31.63	21.49	20.35	17.47	**	+	*	+
NT2RM1000961	4.69	3.03	1.81	5.01	3.8	5.09	4.95	2.93	3.68				
NT2RM1000962	10.02	5.16	7.78	8.82	8.11	7.03	6.17	4.67	6.47				
NT2RM1000973	24.68	15.4	13.27	17.56	15.99	16.81	11.83	13.98	10.68				
NT2RM1000978	0.62	0.04	-0.01	0.17	0.58	0.51	0.69	0.66	1.52				
NT2RM1000982	2.39	1.7	1.71	1.03	0.94	2.7	1.35	1.92	1.56				
NT2RM1000991	4.41	2.48	1.07	2.93	3.33	3.07	1.23	1.71	2.43				
NT2RM1000994	8.78	4.48	6.65	3.77	4.2	8.32	4.28	3.9	4.29				
NT2RM1001002	11.56	5.39	7.09	9.93	9.4	9.55	4.65	6.66	4.14				
NT2RM1001003	9.4	5.64	4.27	5.67	5.91	6.46	6.24	6.75	4.66				
NT2RM1001008	1.85	1.09	0.94	1.76	1.19	2.21	0.79	1.95	1.36				
NT2RM1001011	8.02	5.18	3.04	5.49	6.15	5.88	8.36	7.88	8.53				
NT2RM1001013	2.47	1.58	1.45	1.29	3.7	3.05	2.27	3.51	2.54				
NT2RM1001017	2.77	1.58	1.89	1.79	2.82	2.34	1.35	1.86	1.5				
NT2RM1001018	31.03	16.64	15.26	25.69	26.32	22.96	12.01	17.57	15.08				
NT2RM1001026	5.92	2.62	3.94	6.27	6.63	8.85	2.75	5.72	4.3				
NT2RM1001028	3.4	0.93	2.15	2.01	2.78	3.77	1.36	3.31	2.13				
NT2RM1001043	15.05	7.93	6.39	4.61	4.5	5.16	5.79	4.43	5.13				
NT2RM1001044	4.89	2.09	2.59	3.97	3.59	4.24	2.42	2.42	2.72				
NT2RM1001059	2.09	0.86	1.15	1.37	1.59	1.67	1.46	1.35	0.96				
NT2RM1001063	2.45	1.26	1.65	1.46	2.05	1.8	2.13	2.29	2.06				
NT2RM1001066	1.88	0.18	0.47	1.26	1.05	1.21	0.72	1.03	1.71				
NT2RM1001072	1.32	0.2	0.66	1.3	1.67	2.06	1.25	1.37	0.66				
NT2RM1001074	3.05	0.93	1.31	1.69	2.05	3.12	1.02	1.75	1.85				
NT2RM1001076	1.54	0.37	0.75	0.28	0.39	1.03	0.31	0.72	0.38				
NT2RM1001082	6.04	3.83	2.77	7.68	5.09	7.64	2.86	4.04	3.38				
NT2RM1001085	2.68	0.85	0.53	1.55	1.52	1.92	1.8	2.19	0.8				
NT2RM1001092	7.52	3.6	5.96	8.95	10.4	8.32	6.31	3.61	6.43				
NT2RM1001102	3.26	0.53	1.68	1.38	1.75	2.72	1.2	2.01	1.94				
NT2RM1001103	0.88	0.73	0.28	3.91	4.58	4.4	2.72	2.34	1.98	**	+	**	+
NT2RM1001105	1	0.24	0.43	1.87	1.39	1.31	0.88	1.29	1.26	*	+		
NT2RM1001112	2.67	1.09	1.84	2.3	1.58	2.94	0.99	2.93	1.7				
NT2RM1001115	4.95	1.32	1.99	4.02	5.02	6.62	3.14	4.83	3.48				
NT2RM1001122	8.5	4.16	3.4	8.68	4.04	8.48	4.45	3.73	3.94				
NT2RM1001136	4.05	1.12	0.91	2.5	2.13	2.13	2.47	2.49	2.41				
NT2RM1001139	6.27	3.92	2.62	3.53	3.94	4.14	5.81	5.51	4.63				
NT2RM2000083	2.91	3.18	0.75	4.84	2.4	1.79	5.06	2.26	0.96				
NT2RM2000086	5.44	1.69	3.43	6.16	4.98	7.47	3.88	4.21	4.64				
NT2RM2000090	9.71	5.56	5.39	7.07	8.33	10.49	7.05	5.99	5.68				
NT2RM2000093	2.55	2.71	2.44	3.49	3.87	4.31	1.27	2.57	2.16	**	+		
NT2RM2000096	4.2	1.71	3.04	3.74	3.15	4.87	1.68	3.63	1.98				
NT2RM2000099	14.54	8.15	3.59	5.5	2.42	5.43	3.03	2.67	4.06				
NT2RM2000103	7.04	3.95	5.72	5.91	6.33	6.41	4.47	6.78	5.88				
NT2RM2000106	1.29	2.29	1.74	1.36	3.51	3.21	7.29	2.12	2.85				
NT2RM2000109	8.22	4.26	4.76	1.43	1.14	1.72	2.08	1.91	0.73	*	-	*	-
NT2RM2000112	5.44	2.68	4.48	6.31	4.11	9.84	5.21	4.37	5				
NT2RM2000115	5.58	2.71	2.34	4.26	5.98	6.15	4.54	4.36	4.29				
NT2RM2000118	4.75	4.44	4.18	5.66	3.53	4.65	2.85	2.72	1.51		*	-	
NT2RM2000121	3.3	1.98	1.26	2.86	2.54	1.84	2.16	2.28	2.14				
NT2RM2000124	2.24	1.76	1.1	2.45	4.74	4	2.88	2.71	3.03		*	+	
NT2RM2000127	16.4	9.01	10.98	10.77	15.67	11.6	7.34	7.57	8.37				
NT2RM2000130	3.67	3.12	2.39	2.43	2.62	2.15	1.03	2.04	1.36		*	-	
NT2RM2000133	6.19	3.2	3.93	5.19	4.97	5.78	6.05	5.06	6.76				
NT2RM2000136	21.06	15.5	8.47	21.89	29.21	21.68	13.11	14.37	17.25				
NT2RM2000139	6.65	3.31	3.03	7.38	6.29	6.04	4.13	7.91	5.35				

Table 229

	NT2RM1000691	1.49	0.33	0.72	2.19	3.8	4.38	1.16	2.44	1.23	*	+		
	NT2RM1000698	9.46	4.02	2.95	1.73	2.75	2.69	1.76	3.25	2.56				
5	NT2RM1000699	5.92	1.52	1.15	3.89	3.52	2.89	2.18	3.43	1.91				
	NT2RM1000702	6.62	2.57	3.45	4.5	3.42	3.78	4.09	2.32	3.62				
	NT2RM1000703	17.1	15.01	10.3	10.55	11.96	11.61	8.87	8.94	9.74				
	NT2RM1000704	65.68	42.42	42.04	15.75	17.49	15.71	13.71	12.1	16.78	*	-	*	-
	NT2RM1000725	2.89	1.28	2.86	8.31	19.48	14.98	22.1	28.8	20.7	*	+	**	+
10	NT2RM1000726	2.12	1.3	1.96	2.34	2.21	3.46	1.65	2.75	1.67				
	NT2RM1000731	5.27	2.15	2.93	3.31	4.19	2.99	4.88	3.29	2.95				
	NT2RM1000741	1.93	0.67	1.46	0.89	1.2	1.46	1.17	1.5	1.29				
	NT2RM1000742	23.68	12.81	12.51	8.34	8.53	8.89	7.58	8.47	7.71				
	NT2RM1000744	6.58	2.57	2.31	5.25	4.4	4.66	2.69	3.48	4.72				
15	NT2RM1000746	6.6	3.69	2.39	2.21	4.12	4.39	2.87	3.97	3.11				
	NT2RM1000747	7.04	3.26	3.4	5.08	4.8	5.81	8.95	8.11	9.87		*	+	
	NT2RM1000752	2.53	0.89	1.4	2.34	2.42	2.14	1.42	2.26	1.37				
	NT2RM1000767	7.61	2.5	4.43	7.29	7.21	8.59	10.72	8.37	9.7		*	+	
	NT2RM1000770	5.9	2.04	3.1	5.61	2.94	6.75	3.14	3.37	3.76				
20	NT2RM1000772	2.24	0.1	0.45	1.66	1.02	0.57	0.12	1.61	0.68				
	NT2RM1000779	21.92	14.11	10.14	21.3	25.71	21.61	29.07	21.85	26.85				
	NT2RM1000780	3.49	1.84	0.6	4.74	3.37	4.7	3.33	3.29	1.67				
	NT2RM1000781	0.57	0.24	0.41	1.11	0.76	1.25	0.94	2.16	0.86	*	+		
	NT2RM1000789	3.24	2.46	2.34	3.02	3.98	4.62	2.09	4.84	3.17				
25	NT2RM1000800	7.44	3.44	7.11	8.01	9.85	8.74	6.51	5.53	7.87				
	NT2RM1000802	9.35	5.25	6.84	5.12	5.47	5.85	9.59	9.36	9.75				
	NT2RM1000811	0.9	0.16	0.89	1.36	1.11	1.28	0.91	1.35	0.23				
	NT2RM1000826	26.11	13.59	16.15	23.62	25.62	25.75	12.43	12.08	10.27				
	NT2RM1000829	4.42	3.56	2.62	8.2	6.8	9.18	6.67	6.07	6.37	**	+	**	+
30	NT2RM1000831	96.56	76.65	61.3	78.41	75.7	87	48.08	33.56	47.08		*	-	
	NT2RM1000833	6.27	2.21	1.64	3.09	3.54	4.73	6.47	7.68	4.1				
	NT2RM1000834	4.84	2.51	2.09	5.62	3.9	3.49	3.8	5.68	4.28				
	NT2RM1000841	32.04	19.08	20.07	17.66	18.86	19.57	17.83	9.4	13.52				
	NT2RM1000848	22.37	12.31	11.25	14.54	11.17	13.09	8.36	10.63	15.1				
35	NT2RM1000850	1.25	0.36	0.94	1.01	0.67	1.33	1.5	1.94	1.75		*	+	
	NT2RM1000852	3.74	0.76	1.24	2.68	2.43	2.34	2.39	3.1	1.87				
	NT2RM1000853	1.46	0.57	0.14	1.6	2.87	1.74	1.25	0.52	1.87				
	NT2RM1000855	19.04	8.47	10.06	15.32	18.2	15.69	26.5	18.76	20.5				
	NT2RM1000857	20.9	10.06	10.76	20.92	27.84	24.62	16.83	13.46	17.36				
40	NT2RM1000859	22.68	8.04	9.94	22.93	26.24	26.47	20.88	15.02	18.54				
	NT2RM1000867	15.69	9.11	9.26	15.56	10.14	14.92	15.07	11.26	10.73				
	NT2RM1000874	9.77	5.6	5.03	6.49	6.79	8.79	8.74	7.92	8.94				
	NT2RM1000882	4.01	2.76	2.65	5.69	5.23	6.94	2.13	4.39	2.7	*	+		
	NT2RM1000883	17.32	10.68	13.68	15.2	15.74	17.32	14.61	9.93	20.96				
	NT2RM1000885	31.05	13.08	10.39	19.2	20.71	27.92	20.23	18.36	23.03				
45	NT2RM1000893	3.73	1.65	2.82	3.47	1.63	2.22	4.97	4.49	6.3		*	+	
	NT2RM1000894	14.4	9.62	11.92	7.88	9.3	10.29	9.51	9.36	13.18				
	NT2RM1000896	2.53	0.85	1.96	3.01	2.71	4.11	3.76	3.77	6.2		*	+	
	NT2RM1000899	1.45	0.26	1.26	1.48	1.2	1.14	1.07	1.69	0.72				
	NT2RM1000905	55.04	22.33	30.63	36.24	41.24	41.41	17.87	22.74	23.3				
50	NT2RM1000910	7.05	2.93	6.34	6.29	7.41	5.83	7.31	6.05	5.79				
	NT2RM1000914	8.32	4.94	2.32	6.53	12.83	8.37	4.34	8.86	6.23				
	NT2RM1000919	4.65	2.11	2.49	5.45	3.02	4.5	2.58	2.81	4.74				
	NT2RM1000921	2.3	0.73	0.47	1.57	1	2.01	1.98	1.39	1.88				
	NT2RM1000922	7.7	4.51	3.3	6.07	5.41	6.35	3.4	3.21	3.38				
55	NT2RM1000924	3.33	1.7	1.15	2.35	2.35	2.87	1.24	2.25	1.77				
	NT2RM1000927	3.83	1.15	1.76	5.16	2.77	6.27	2.3	3.24	2.14				
	NT2RM1000951	8.45	4.91	4.93	9.07	6.69	6.96	5.29	7.06	3.5				

Table 228

	NT2RM1000271	0.75	0.2	0.04	1.21	0.35	0.58	0.94	0.84	0.49				
	NT2RM1000272	54.56	36.55	40.59	39.42	48.05	51.89	35.16	41.56	36.18				
5	NT2RM1000273	25.51	11.38	15.12	14.18	12.87	14.49	8.99	9.27	12				
	NT2RM1000274	58.21	39.03	46.94	45.24	44.74	49.05	21.9	22.39	26.39		*	-	
	NT2RM1000280	3.79	2.05	1.14	3.65	3.57	2.6	4.36	3.9	4.03				
	NT2RM1000295	1.04	0.33	0.49	1.43	1.42	1.12	1.49	1.59	1.89	*	+	*	+
	NT2RM1000300	3.37	1.19	1.93	2.35	3.27	3.66	2.84	2	3				
10	NT2RM1000304	119.7	75.04	105.1	129.6	102.4	124.9	50.36	59.48	58.8		*	-	
	NT2RM1000314	14.79	10.41	9.09	12.21	10.45	12.98	11.38	9.76	12.93				
	NT2RM1000318	24.15	19.1	20.62	18.95	25.93	22.36	13.38	12.74	12.13		**	-	
	NT2RM1000335	2.7	1.54	1.86	2.64	0.98	2.51	2.11	1.75	0.87				
	NT2RM1000341	1.86	1.47	0.19	1.35	0.97	1.03	1.64	1.09	1.69				
15	NT2RM1000350	12.53	6.61	5.41	9.68	8.63	6.11	10.39	8.69	12.6				
	NT2RM1000354	1.42	1.08	1.09	1.11	0.94	2.05	1.14	0.93	0.85				
	NT2RM1000355	24.12	12.19	10.53	22.94	22.89	22.53	40.93	26.81	41.82		*	+	
	NT2RM1000361	3.67	1.47	2.35	2.55	2.08	2.7	1.88	1.68	2.1				
	NT2RM1000365	1.06	0.28	0.15	0.8	0.83	1.19	0.3	0.84	1.1				
20	NT2RM1000372	20.32	11.77	14.09	12.5	15.42	19.07	11.35	13.11	12.12				
	NT2RM1000377	4.71	2.13	0.97	3.33	3.33	3.84	3.13	2.47	2.45				
	NT2RM1000388	4.08	1.38	1.89	2.94	1.24	2.04	2.06	1.76	3.15				
	NT2RM1000394	1.97	0.69	0.13	1.46	1.54	2.03	0.91	0.83	1.86				
	NT2RM1000399	1.06	0.34	0.04	1.59	1.17	1.07	1.01	1.52	1.09				
25	NT2RM1000407	3.28	1.69	1.8	2.92	2.58	2.42	3.74	2.39	2.69				
	NT2RM1000421	1.21	0.17	0.31	0.84	0.59	1.24	0.64	0.87	1.2				
	NT2RM1000422	184.9	121.2	142.5	178.6	203	174.3	67.17	77.47	67.99		*	-	
	NT2RM1000430	2.25	0.23	1.58	0.73	1.22	1.54	1.8	1.12	1.6				
	NT2RM1000462	11.14	6.84	5.58	14.5	17.82	8.39	4.89	8.25	6.36				
30	NT2RM1000499	5.37	2.3	2.51	3.94	5.62	7.36	4.89	3.83	3.47				
	NT2RM1000512	22.47	26.43	20.07	26.5	33.66	27.9	17.58	19.86	18.1				
	NT2RM1000519	29.78	19.56	14.02	7.45	11.19	11.75	14.89	14.37	13.43				
	NT2RM1000527	18.16	11.14	6.22	5.88	7.16	7	1.98	1.37	2.55		*	-	
	NT2RM1000539	12.49	8.93	7.21	6.18	6.43	8.69	2.33	4.94	2.74		*	-	
35	NT2RM1000542	5.88	1.72	2.37	3.23	3.3	5.23	2.07	2.93	2.21				
	NT2RM1000553	3.65	0.83	1.64	1.16	1.39	3.69	1.46	2.07	1.37				
	NT2RM1000555	54.21	28.45	27.23	49.44	36.73	39.14	24.87	25.09	25.78				
	NT2RM1000558	5.67	1.77	2.83	4.02	2.67	3.58	2.91	2.6	1.85				
	NT2RM1000563	5.22	2.56	1.89	2.43	2.32	3.96	2.78	2.56	3.17				
	NT2RM1000566	7.28	3.71	3.24	1.61	1.5	1.27	1.81	1.72	3.16				
40	NT2RM1000570	26.49	17.4	16.59	16.76	14.37	17.79	32.95	44.77	33.54		*	+	
	NT2RM1000571	6.81	1.94	3.76	2.38	2.48	3.22	3.14	4.7	3.91				
	NT2RM1000574	1.29	0.74	0.74	1.47	2.46	0.57	1.31	2.11	1.66				
	NT2RM1000580	1.69	0.26	0.99	1.9	0.77	2.4	1.57	1.93	1.37				
	NT2RM1000620	10.67	5.15	5.67	13.49	19.91	14.9	8.69	7.05	7.31	*	+		
45	NT2RM1000623	1.16	0.68	-0.02	1.17	0.81	0.94	0.79	1.05	0.97				
	NT2RM1000630	2.05	1.24	0.77	1.67	2.19	1.83	1.87	1.47	1.67				
	NT2RM1000633	27.41	17.8	20.59	31.5	32.02	35.16	25.03	10.45	15.13	*	+		
	NT2RM1000634	2.52	1	0.44	1.48	1.48	2.34	1.07	2.17	0.81				
	NT2RM1000642	6.47	2.65	3.78	2.95	2.59	5.44	3.92	6.2	4.71				
50	NT2RM1000647	37.58	16.74	23.8	20.92	20.8	27.77	21.56	20.3	24.27				
	NT2RM1000648	2.04	0.41	0.84	1.58	1.79	2.24	1.08	2.71	1.07				
	NT2RM1000650	3.85	1.26	1.26	3.06	2.28	2.37	2.52	2.44	1.64				
	NT2RM1000661	6.75	4.13	3.51	3.45	3.05	3.31	3.37	4.05	2.46				
	NT2RM1000666	25.38	12.8	15.49	0.79	0.92	0.69	0.89	0.98	0.83	*	-	*	-
55	NT2RM1000669	3.69	1.54	2.15	2.54	2.09	3.44	2.17	1.76	1.75				
	NT2RM1000672	18.91	9.34	13.85	40.77	46.8	48.09	11.37	11.65	12.71	**	+		
	NT2RM1000681	7.08	2.25	3.13	16.21	18.15	17.39	29.47	23.33	30.9	**	+	**	+

Table 227

	MAMMA1003140	1.69	0.85	0.6	1.74	1.79	2.62	1.07	1.84	1.32				
	MAMMA1003146	3.17	0.64	1.39	1.41	2.11	2.74	2.14	2.47	2.39				
5	MAMMA1003150	14.6	5.76	6.23	12.63	11.24	8.45	5.72	10.32	7.8				
	MAMMA1003154	8.12	5.17	3.61	5.99	6.38	3.91	3.93	4.78	3.8				
	MAMMA1003155	3.73	2.43	2.74	2.68	3.2	4.47	4.25	3.56	2.96				
	MAMMA1003157	3.72	2.17	1.5	8.43	9.53	5.52	5.81	5.42	4.11	*	+	*	+
	MAMMA1003163	3.24	2.63	2.53	2.86	3.42	4.51	2.32	3.21	3.84				
10	MAMMA1003164	4.04	1.62	1.78	2.36	3.89	3.12	1.98	3.3	1.9				
	MAMMA1003166	2.64	0.97	1.34	1.14	2.03	2.6	0.94	1.46	0.67				
	NB9N31000010	14.76	5.71	8.03	2.59	3.65	2.88	2.38	3.3	2.11				
	NB9N31000016	7.03	5.06	4.31	4.14	3.19	3.67	2.48	3.3	3.54				
	NB9N31000043	6.43	3.37	2.66	3.6	4.63	3.3	4.03	4.8	3.85				
15	NB9N31000045	19.15	14.02	9.92	7.25	11.2	10.47	9.72	9.74	10.85				
	NB9N31000054	6.46	2.26	2.68	6.57	6.74	6.4	6.39	5.01	4.4				
	NB9N31000076	2.64	1.86	1.23	4.27	5.28	5.06	3.51	3.29	3.06	**	+	*	+
	NB9N31000086	3.3	1.41	1.24	4.91	5.73	5.78	4.03	3.25	4.37	**	+		
	NT2RM1000001	3.65	2.34	1.78	2.42	3.06	4.27	1.46	2.56	2.65				
20	NT2RM1000018	18.02	4.88	9.18	11.8	18.97	15.96	10.32	8.58	7.34				
	NT2RM1000032	2.53	0.99	1.56	3.18	2.12	2.58	1.32	2.6	2.8				
	NT2RM1000035	11.4	5.02	6.42	9.17	9.42	10.51	8.5	7.07	7.86				
	NT2RM1000037	13.15	8.99	9.27	10.68	10.22	12.08	8.43	7.97	9.91				
	NT2RM1000039	11.18	9.88	11.7	14.16	13.27	16.95	11.97	10.55	15.86	*	+		
25	NT2RM1000042	80.13	61.43	48.95	80.07	94.16	101.1	34.69	35.38	37.43		*	-	
	NT2RM1000053	1.63	0.44	0.19	1.9	1.2	1.06	0.56	1.65	0.56				
	NT2RM1000059	10.72	6.4	6.93	10.31	13.85	13	8.96	10.38	10.03				
	NT2RM1000062	2	0.27	0.62	1.05	1.09	1.16	1.09	1.18	1.04				
	NT2RM1000065	113.3	91.26	69.94	64.48	58.5	52.5	34.11	33.99	50.67		*	-	
30	NT2RM1000066	35.22	18.22	21.68	21.61	23.29	23.48	22.94	24.27	17.75				
	NT2RM1000071	63.91	66.46	45.7	62.4	99.26	85.6	34.56	28.87	36.37		*	-	
	NT2RM1000080	3.9	1.47	1.12	2.18	2.14	2.55	1.54	2.09	2.44				
	NT2RM1000086	19.75	10.02	12.84	15.85	21.11	21.57	16.5	12.7	16.82				
	NT2RM1000092	3.84	1.47	1.22	4.35	3.45	3.58	5.38	4.65	2.8				
	NT2RM1000118	0.16	0.1	0.44	0.44	0.48	0.43	0.45	1.71	0.2				
35	NT2RM1000119	1.47	0.16	1.14	1.49	1.8	1.27	0.45	3.87	1.63				
	NT2RM1000121	3.95	2.18	1.02	2.75	2.63	2.42	2.12	2.47	2.71				
	NT2RM1000122	20.69	10.42	10.67	11.66	9.11	15.06	12.71	8.89	10.81				
	NT2RM1000127	3.09	0.8	1.57	1.55	1.35	2.79	1.74	2.29	1.61				
	NT2RM1000131	1.39	0.57	0.54	0.93	0.82	1.7	1.32	1.99	1.76				
40	NT2RM1000132	3.41	2.17	2.19	3.36	2.6	3.36	3.07	3.21	1.8				
	NT2RM1000153	2.4	1.2	1	2.3	1.9	1.72	2.33	2.75	1.99				
	NT2RM1000184	12.46	9.34	11.07	12.61	11.31	13.35	27.02	24.07	25.86		**	+	
	NT2RM1000186	0.96	0.05	1.17	1.92	0.66	0.6	1.01	1.84	0.71				
	NT2RM1000187	7.97	7.07	3.88	7.69	10.3	6.3	4.37	5.12	5.93				
45	NT2RM1000199	2.43	1.17	0.94	2.23	1.56	2.06	2.22	2.21	0.97				
	NT2RM1000213	4.77	2.05	1.72	5.31	3.68	5.55	3.01	2.88	2.04				
	NT2RM1000215	22.27	12.67	13.12	16.61	13.19	18.25	21.54	17.2	19.32				
	NT2RM1000218	4.96	1.49	2.25	6.26	5.56	5.79	6.91	7.15	6.31		*	+	
	NT2RM1000224	14.47	7.85	6.71	14.79	9.45	14.6	5.58	5.7	6.61				
50	NT2RM1000236	11.3	7.18	4.01	4.18	3.11	5.39	11.74	17.39	13.83				
	NT2RM1000242	-0.07	0.1	-0.14	0.21	0.11	0.96	0.08	1.21	-0.15				
	NT2RM1000244	3.77	1.77	0.73	1.27	1.58	1.27	0.95	0.89	0.64				
	NT2RM1000252	31.79	17.18	15.85	26.05	28.99	28.97	19.44	15.24	17.84				
	NT2RM1000256	20.24	12.26	8.91	9.14	16.96	11.49	13.69	10.22	13.24				
	NT2RM1000257	16.34	9.13	9.74	4.83	6.53	7.09	5.1	3.9	3.96		*	-	
55	NT2RM1000260	32.33	14.76	16.72	33.82	31.04	33.35	23.69	23.35	23.61				
	NT2RM1000269	12.22	8.71	9.44	5.25	3.16	4.72	1.74	2.23	1.16	**	-	**	-

Table 226

	MAMMA1002895	1.52	1.02	0.66	3.67	2.82	2.63	1.68	3.27	1.67	**	+		
	MAMMA1002898	5.3	1.67	2.43	5.04	3.66	3.54	3.19	4.2	4.28				
5	MAMMA1002905	7.3	4.24	4.9	4.36	3.31	5.5	4.49	4.07	7.6				
	MAMMA1002906	7.09	3.55	2.11	4.13	4.15	4.17	3.6	4.08	4.37				
	MAMMA1002908	5.1	3.63	2.55	7.12	10.01	7.24	3.97	3.94	6.08	*	+		
	MAMMA1002909	11.19	2.36	4.9	18.65	20.5	19.49	11.96	9.14	7.19	**	+		
	MAMMA1002918	8.8	4.28	4.36	7.71	4.97	6.64	4.29	3.86	3.85				
10	MAMMA1002925	3.35	2.63	1.48	9.46	7.99	8.84	13.12	8.46	14.83	**	+	**	+
	MAMMA1002926	7.82	4.53	3.55	10.54	8.94	10.54	4.02	3.98	2.94	*	+		
	MAMMA1002930	4.28	1.73	3.17	5.74	5.95	7.07	4.01	5.04	2.57	*	+		
	MAMMA1002937	5.96	2.45	3.44	4.74	4.53	5.73	3.19	3.43	4.76				
	MAMMA1002938	3.7	2.19	0.47	2.73	4.56	4.15	4.37	4.59	4.01				
15	MAMMA1002941	1.15	1.12	0.39	3.44	2.75	4.14	1.85	1.74	2.91	**	+	*	+
	MAMMA1002947	6.2	1.75	2	3.69	4.63	4.41	4	2.74	2.53				
	MAMMA1002964	3.13	0.8	1.6	3.89	5.54	5.2	2.56	3.32	2.95	*	+		
	MAMMA1002967	2.77	0.81	0.72	2.65	3.25	3.1	2.15	2.36	2				
	MAMMA1002970	10.68	5	6.77	15.62	18.38	19.77	9.12	10.6	10.22	**	+		
20	MAMMA1002971	5.36	1.91	2.72	5.34	4.3	4.54	3.53	5.4	3.71				
	MAMMA1002972	3.58	1.23	1.8	5.51	3.48	3.8	2.78	4.51	3.78				
	MAMMA1002973	3.05	2.45	2.19	5.84	7.86	5.49	3.04	3.4	3.84	**	+		
	MAMMA1002979	49.45	21.28	20.21	54.78	50.04	57.56	26.52	29.51	38.14				
	MAMMA1002982	1.17	0.84	0.21	1.07	1.04	1.44	0.75	0.85	2.52				
25	MAMMA1002987	2.51	2.1	1.94	4.65	4.24	4.32	2.66	3.22	2.69	**	+		
	MAMMA1003003	6.44	2.24	3.39	6.63	8.14	8.81	3.38	3.94	4.55				
	MAMMA1003004	2.44	1.12	1.78	4.34	4.64	5.27	2.45	2.33	3.36	**	+		
	MAMMA1003007	3	0.97	0.37	1.72	3.13	2.66	1.67	2.02	2.34				
	MAMMA1003011	6.89	3.86	2.58	10.11	6.23	6.02	5.56	4.68	6.89				
30	MAMMA1003013	4.71	2.5	3.6	5.96	2.57	4.98	4.47	2.47	4.04				
	MAMMA1003015	3.11	1.7	0.83	3.85	3.23	4.39	2.92	3.35	3.6				
	MAMMA1003019	1.94	0.48	0.77	1.44	1.99	1	1.47	1.37	1.39				
	MAMMA1003020	4.98	3.11	2.83	4.85	4.06	4.94	3.36	4.67	2.34				
	MAMMA1003026	2.22	1.04	1.33	2.17	1.21	1.23	1.15	1.94	1.66				
35	MAMMA1003031	10.83	4.3	5.89	8.39	13.69	12.78	6.3	8.07	8.55				
	MAMMA1003033	4.26	3.18	1.65	3.05	5.95	7.17	2.79	4.73	3.1				
	MAMMA1003035	9.17	3.04	2.57	6.09	5.43	4.4	3.27	3.33	2.99				
	MAMMA1003039	2.73	0.66	0.77	3.23	4.07	2.57	2.03	1.92	2.63				
	MAMMA1003040	5.92	4.5	4.4	12.47	14.15	15.98	6	7.82	5.59	**	+		
	MAMMA1003044	5.54	1.89	2.06	8.57	6.1	5.51	3.66	3.75	3.73				
40	MAMMA1003047	24.49	9.27	14.52	16.47	16.89	16.3	13.85	12.65	14.22				
	MAMMA1003049	1.66	0.7	0.16	1.59	1.6	1.36	1.06	0.97	1.99				
	MAMMA1003055	3.44	1.83	1.31	3.88	3.78	5.3	1.65	3.16	2.91				
	MAMMA1003056	3.11	0.29	1.13	1.54	2.14	2.78	1.67	3.29	1.4				
	MAMMA1003057	4.22	3.06	2.41	5.23	4.85	4.4	3.28	3.47	3.84				
45	MAMMA1003066	4.41	2.68	2.13	7.59	8.47	7.26	3.45	3.84	3.94	**	+		
	MAMMA1003075	2.52	1.24	0.49	2.49	1.99	2.02	1.98	1.74	1.75				
	MAMMA1003089	3.39	2.37	1.55	7.01	9.09	5.24	3.86	3.79	4.04	*	+		
	MAMMA1003092	2.28	2.1	0.75	1.76	2.8	2.59	1.29	2.14	0.99				
	MAMMA1003095	3.31	3.21	2.49	5.68	6.41	6.17	3.79	3.05	2.04	**	+		
50	MAMMA1003099	4.62	1.71	1.38	5.27	3.36	5.17	3.64	4.12	3.25				
	MAMMA1003102	4.98	1.87	1.62	3.02	1.85	3.96	2.26	3.51	2.66				
	MAMMA1003104	3.42	0.58	0.51	3.71	2.62	2.21	2	1.93	0.79				
	MAMMA1003113	7.31	2.8	2.6	2.96	4.59	4.44	3.67	4.2	3.98				
	MAMMA1003126	5.27	3.19	2.59	5.46	4.59	6.5	4.92	4.89	5.03				
55	MAMMA1003127	3.2	0.92	0.9	2.49	3.46	2.32	2.81	3.43	2.3				
	MAMMA1003131	14.8	5.77	8.64	6.66	11.84	10.58	7.76	9.54	7.3				
	MAMMA1003135	2.29	0.95	1.03	2.13	1.48	2.22	0.96	2.28	1.38				

Table 225

	MAMMA1002701	5.66	2.9	4.33	9.27	7.16	8.59	4.61	5.08	5.08	*	+		
	MAMMA1002708	7.94	5.73	7.17	9.47	9.6	11.7	5.3	7.78	6.06	*	+		
5	MAMMA1002711	5.14	1.55	3.02	5.08	5.35	9.25	4.88	5.17	3.67				
	MAMMA1002712	8.23	3.4	3.83	5.92	5.37	4.49	4.33	4.65	3.86				
	MAMMA1002716	3.03	1.15	1.75	3.45	3.66	6.18	3.63	4.99	6.27		*	+	
	MAMMA1002721	5.09	3.43	2.39	8.57	10.12	9.06	4.73	4.05	4.78	**	+		
	MAMMA1002723	3.9	1.75	1.64	3.74	4.55	4.64	2.71	2.75	3.13				
10	MAMMA1002727	1.94	0.37	0.28	1.65	1.68	1.6	1.31	1.6	1.09				
	MAMMA1002728	18.85	12.15	13.58	19.57	15.85	19.98	10.65	11.63	8.96				
	MAMMA1002742	24.64	11.73	11.42	17.86	18.78	18.95	12.46	17.75	16.29				
	MAMMA1002743	3.32	1.38	1.48	2.64	3.77	2.84	1.3	3.55	2.08				
	MAMMA1002744	5	2.18	1.83	8.37	6.2	7.98	3.63	3.32	2.37	*	+		
15	MAMMA1002746	2.51	0.63	0.79	1.49	2.16	1.83	2.14	1.51	0.81				
	MAMMA1002748	3.99	1.96	1.48	3.96	2.53	5.35	2.11	2.64	2.6				
	MAMMA1002754	3.27	1.38	1.23	3.72	4.67	3.51	3.5	2.37	3.36				
	MAMMA1002758	1.75	1.23	0.68	1.23	1.77	1.88	1.75	1.78	0.81				
	MAMMA1002762	15.53	11.07	116.89	14.23	17.23	16.31	8.35	12.66	9.99				
20	MAMMA1002764	6.2	2.6	2.93	8.75	9.77	8.81	4.73	4.74	4.79	*	+		
	MAMMA1002765	4.28	1.57	1.43	2.94	4.93	4.38	2.62	3.87	2.62				
	MAMMA1002769	1.56	0.46	0.63	2.76	2.64	1.76	3.07	2.6	2.53	*	+	**	+
	MAMMA1002771	7.14	1.91	2.56	3.71	2.39	3.56	2.38	4.39	2.84				
	MAMMA1002775	8.17	3.51	3.32	3.63	6.17	5.65	3.96	3.51	3				
25	MAMMA1002780	4.25	0.67	1.1	3.25	4.36	3.86	1.61	2.45	1.84				
	MAMMA1002782	3.73	1.77	1.35	3.47	4.14	4.44	2.59	3.58	3.12				
	MAMMA1002795	1.54	0.63	0.41	1.27	1.55	2.07	1.2	2.31	1.82				
	MAMMA1002796	5.26	2.04	2.88	2.31	3.68	4.71	3.08	4.01	2.78				
	MAMMA1002805	1.95	1.42	2.03	2.66	2.54	2.92	1.33	2.31	1.29	*	+		
	MAMMA1002806	7.18	3.13	2.76	7.9	8.06	6.82	4.84	4.21	4.71				
30	MAMMA1002807	5.28	1.74	0.98	3.68	4.66	5.86	3.42	3.27	3.02				
	MAMMA1002814	3.87	2.51	3.12	7.45	7.16	7.74	4.16	4.93	4.92	**	+	*	+
	MAMMA1002817	1.7	0.51	0.6	1.42	1.13	1.4	0.99	1.61	0.6				
	MAMMA1002820	1.34	1.92	0.86	2.57	2.4	3.83	1.38	1.74	1.69	*	+		
	MAMMA1002830	27.11	10.85	16.25	30.04	35.58	32.67	18.44	20.75	20.74	*	+		
35	MAMMA1002833	6.78	4.02	4.05	10.31	9.78	13.03	4.43	6.24	5.25	*	+		
	MAMMA1002835	3.11	0.73	1.29	2.37	4.3	3.68	1.9	2.74	1.11				
	MAMMA1002838	5.08	1.94	1.5	7.62	5.02	5.3	2.99	3.7	3.52				
	MAMMA1002842	6.45	2.71	2.75	6.39	9.1	5.17	5.25	5.53	5.55				
	MAMMA1002843	4.18	1.22	2.78	4.36	3.92	4.27	2.84	3.41	2.54				
40	MAMMA1002844	15.29	8.97	10.98	13.02	14.25	13.61	12.26	13.86	18.37				
	MAMMA1002845	0.94	0.26	0.38	2.62	1.75	2.18	15.33	12.73	11.67	**	+	**	+
	MAMMA1002857	92.97	61.45	71.01	93.18	91.48	102.4	49.65	49.13	49.57				
	MAMMA1002858	270.3	178.2	193.7	198.5	285	325.3	136.6	154.1	144.4				
	MAMMA1002863	6.79	3.17	3.17	4.69	5.56	4.89	3.85	6.3	4.27				
45	MAMMA1002868	5.34	2.46	2.35	7.72	6.47	7.85	3.3	3.69	4.4	*	+		
	MAMMA1002869	6.13	2.1	3.45	4.16	4.01	5.84	3.15	3.68	3.35				
	MAMMA1002871	0.97	0.66	0.13	2.7	2.82	2.55	1.36	2.18	2.35	**	+	*	+
	MAMMA1002875	4.77	2.06	2.53	6.78	7.19	6.9	3.55	3.8	4.77	*	+		
	MAMMA1002879	3.84	2.9	2.39	3.98	5.13	4.33	4.2	4.17	4.62		*	+	
	MAMMA1002880	3.28	1.24	0.99	2.01	1.85	1.9	3.12	3.06	1.4				
50	MAMMA1002881	5.17	2.92	2.09	6.15	9.22	4.65	3.67	4.57	4.68				
	MAMMA1002885	5.25	2.85	2.52	4.49	4.87	6.69	2.72	4.39	2.98				
	MAMMA1002886	6.24	3.43	2.66	5.52	4.58	6.71	3.64	3.49	2.72				
	MAMMA1002887	3.89	0.95	1	1.97	1.75	1.77	2.34	1.93	1.82				
	MAMMA1002890	5.13	2.67	3.05	8.31	4.7	8.72	5.27	5.4	5.01				
55	MAMMA1002892	5.88	3.48	2.47	7.32	8.24	6.42	4.17	5.23	4.57	*	+		
	MAMMA1002893	8.86	9.67	8.59	8.18	9.34	9.39	5.69	3.91	5.29		**	-	

Table 224

	MAMMA1002466	7.61	3.8	3.03	7.05	8.64	7.32	9.99	8.37	11.38			*	+
	MAMMA1002470	5.61	2.03	2.45	2.62	3.83	4.24	2.19	2.79	3.07				
5	MAMMA1002475	2.73	2.58	1.69	4.8	5.81	4.75	1.5	3.35	3.39	**	+		
	MAMMA1002480	1.82	0.76	1.1	1.61	2.6	1.72	0.67	1.56	1.72				
	MAMMA1002485	11.15	6.59	4.25	5.55	8.76	7.85	6.2	6.28	8.64				
	MAMMA1002494	6.22	5.16	3	7.41	9.6	7.67	4.89	3.44	6.03	*	+		
	MAMMA1002498	5.71	3.03	1.34	3.92	2.98	3.69	2.66	2.39	3.29				
10	MAMMA1002524	7.17	3.31	2.26	5.6	4.65	6.85	3.63	4.86	5.05				
	MAMMA1002530	5.79	3.23	2.55	4.12	8.81	3.19	5.21	4.47	5.09				
	MAMMA1002538	4.01	3.96	2.85	3.37	4.2	2.1	2.88	2.7	3.45				
	MAMMA1002545	8.19	4.19	5.05	10.66	9.93	10.97	4.47	4.9	6.19	*	+		
	MAMMA1002554	4	1.52	3.49	3.57	3.68	3.97	1.82	2.91	3.1				
15	MAMMA1002556	9.93	4.82	2.86	7.06	11.34	10.05	5.76	5.07	5.23				
	MAMMA1002561	10.06	3.9	4.44	12.05	12.4	15.05	9.97	6.01	8.09	*	+		
	MAMMA1002565	4.89	4.2	3.26	4.07	7.56	4.55	3.68	2.91	4.58				
	MAMMA1002566	4	2.15	0.94	5.93	2.4	2.55	2.16	2.54	3.99				
	MAMMA1002571	7.22	3.36	3.15	5.32	6.04	4.33	4.11	4.2	3.94				
20	MAMMA1002573	11.2	4.78	6.52	15.53	15.17	13.55	7.02	8.07	9.44	*	+		
	MAMMA1002576	6.01	1.71	4.22	10.04	10.33	6.3	4	6.04	6.94				
	MAMMA1002584	11.01	7.77	8.72	19.33	19.85	20.62	8.27	12.03	12.19	**	+		
	MAMMA1002585	7.85	4.99	2.28	4.43	8.97	3.79	4.59	2.67	4.69				
	MAMMA1002586	4.6	2.19	2.47	3.71	4.21	5.32	2.84	2.51	4.3				
25	MAMMA1002589	4.94	2.94	1.69	6.3	6.89	4.51	3.93	3.36	4.69				
	MAMMA1002590	10.71	5.82	7.42	10.33	15.26	8.36	9.91	9.3	15.5				
	MAMMA1002593	7.21	1.7	2.9	10.38	6.09	7.62	3.83	4.23	4.78				
	MAMMA1002597	5.27	4.72	2.89	5.79	7.99	6.52	3.32	4.98	3.89				
	MAMMA1002598	28.18	14.66	17.3	23.76	26.47	26.12	9.35	11.37	10.26				
30	MAMMA1002603	3.82	2.48	2.87	6.45	7.78	6.16	3.06	4.45	5.16	**	+		
	MAMMA1002612	18.88	8.49	7.35	14.76	23.79	19.09	13.04	8.06	15.93				
	MAMMA1002617	20.5	11.92	10.78	21.62	26.8	21.46	18.22	10.24	15.46				
	MAMMA1002618	8.07	5.37	4.36	5.18	5.81	5.01	3.29	4.53	3.87				
	MAMMA1002619	2.75	1.98	1.32	3.42	3.69	3.38	3.52	2.56	2.73	*	+		
35	MAMMA1002622	4.65	2.19	2.57	6.98	7.16	7	3.88	4.47	5.21	**	+		
	MAMMA1002623	3.7	4.09	2.66	8.45	8.43	10.17	4.49	5.06	6.96	**	+		
	MAMMA1002625	1.31	0.77	1.1	4.74	4.02	3.9	1.84	3.63	1.92	**	+		
	MAMMA1002627	0.15	0.77	0.52	0.63	0.61	1.31	0.61	0.89	0.31				
	MAMMA1002629	5	1.49	4.04	8.25	13.1	6.87	3.59	5.41	7.41				
	MAMMA1002631	3.02	0.94	0.62	3.54	2.28	2	1.53	1.32	2.73				
40	MAMMA1002633	8.62	2.1	5.7	4.72	6.74	7.92	3.72	4.69	4.2				
	MAMMA1002636	3.59	1.19	1.71	4.59	3.63	5.19	2.99	3.81	3.18				
	MAMMA1002637	1.74	1.17	1.01	2.51	1.67	1.58	1.65	2.79	2.2				
	MAMMA1002646	5.71	2.6	2.44	4.61	4.24	4.68	2.72	3.67	2.73				
	MAMMA1002648	9.62	6.84	5.82	8.64	14.71	12.83	6.98	7.83	7.07				
45	MAMMA1002650	0.72	0.4	0.49	1.46	0.42	1.02	0.69	0.35	0.84				
	MAMMA1002652	6.32	1.69	4.33	6.84	5.22	9.05	3.61	3.81	5.06				
	MAMMA1002655	6.13	2.3	1.98	3.61	1.81	5.05	3.34	3.19	3.44				
	MAMMA1002662	5.15	2.31	2.11	6.95	6.87	5.4	4.01	4.49	5.25	*	+		
	MAMMA1002665	11.8	6.1	10.13	10.87	17.41	15.49	7.23	8.06	7.62				
	MAMMA1002671	7.41	2.14	3.42	5.62	4.48	5.33	3.61	3.41	3.76				
50	MAMMA1002673	7.4	3.46	4.23	7.31	8.7	9.27	5.9	6.54	4.84				
	MAMMA1002684	9.53	3.22	5.59	4.24	7.51	8.57	6.73	6.88	7.64				
	MAMMA1002685	3.8	1.88	0.7	2.75	4.35	3.69	1.82	1.26	1				
	MAMMA1002692	7.2	4.36	3.76	8	7.57	6.47	4.09	3.19	4.9				
	MAMMA1002693	8.11	3.16	4.22	9.2	3.75	8.99	4.65	5.78	5.13				
55	MAMMA1002698	5.29	1.74	2.15	6.64	6.43	7.76	3.35	3.54	3.9	*	+		
	MAMMA1002699	2.23	0.61	0.97	1.33	2.22	1.52	1.64	1.71	1.92				

Table 223

	MAMMA1002267	5.13	1.56	2.1	4.1	8	6.58	5.59	7.23	7.33		*	+
	MAMMA1002268	4.34	3.93	2.18	3.97	3.15	4.33	1.93	3.77	3.06			
5	MAMMA1002269	3.53	2.77	0.37	2.27	1.57	2.25	1.64	1.13	1.9			
	MAMMA1002282	3.17	4.02	1.28	2.38	4.52	4.47	2.52	2.77	2.84			
	MAMMA1002292	8	3.86	4.57	6.11	4.23	6.12	4.47	3.55	4.28			
	MAMMA1002293	13.94	6.19	6.42	18.8	17.8	21.12	10.21	8.07	15.59	*	+	
	MAMMA1002294	6.97	4.11	3.04	6.45	7.32	6.27	5.03	5.25	5.73			
10	MAMMA1002297	5.17	2.14	2.44	5.18	5.03	6.05	4.2	2.91	4.33			
	MAMMA1002298	5.95	2.63	2	5.32	4.87	5.66	3.33	3.41	4.16			
	MAMMA1002299	3.71	2.19	2.17	3.02	3.23	3.18	3.21	2.61	2.25			
	MAMMA1002308	4.09	3.82	1.96	6.63	7.73	3.7	2.44	2.99	3.59			
	MAMMA1002310	24.32	15.32	19.7	26.21	29.99	31.31	20.38	19.58	18.88	*	+	
15	MAMMA1002311	10.38	6.89	2.86	14.02	13.82	13.05	10.49	6.04	10.98	*	+	
	MAMMA1002312	7.11	4.07	0.96	3.66	5.77	5.39	2.87	2.07	3.97			
	MAMMA1002317	5.37	4.98	2.41	6.38	13.31	8.87	4.49	3.92	7.76			
	MAMMA1002319	8.07	2.35	5.23	7.19	7.92	8.72	5.3	5.48	6.56			
	MAMMA1002322	6.31	4.11	5.15	10.22	11.41	12.06	4.9	7.5	6	**	+	
20	MAMMA1002329	4.15	2.37	1.67	2.9	3.82	5.04	2.2	3.87	3.47			
	MAMMA1002332	4.13	2.74	1.9	3.61	6.19	6.87	2.13	3.26	3.02			
	MAMMA1002333	7.26	4	2.1	6.05	5.74	3.04	3.25	4.13	4.42			
	MAMMA1002335	10.93	3.6	4.03	10.38	8	8.37	5.57	5.29	6.32			
	MAMMA1002339	7.73	3.96	3.73	8.81	10.04	9.53	3.71	3.46	7.48	*	+	
25	MAMMA1002347	6.93	4.17	2.03	4.83	7.45	7.07	4.3	4.21	4.94			
	MAMMA1002351	3.84	5.05	2.4	3.45	5.38	4.65	4.23	5.29	5.91			
	MAMMA1002352	5.21	4	2.14	4.04	3.97	4.72	2.11	1.72	2.04			
	MAMMA1002353	9.22	7.52	2.31	5.95	8.94	7.55	4.37	4.54	4.03			
	MAMMA1002355	5.34	3.25	2.3	4.76	5.27	7.77	2.43	4.79	2.85			
	MAMMA1002356	3.57	2.35	1.19	3.19	4.03	4.8	2.05	2.5	2.26			
30	MAMMA1002359	13.77	9.98	8.17	18.6	20.01	21.01	10.51	7.95	8.5	**	+	
	MAMMA1002360	4.19	2.61	1.63	3.14	2.98	2.4	3	1.64	2.41			
	MAMMA1002361	6.53	2.69	2.54	6.26	7.25	5.96	4.09	4.49	5.12			
	MAMMA1002362	3.93	2.21	1.89	3.56	5.61	4.11	4.72	2.96	3.12			
	MAMMA1002367	6.65	2.94	3.45	4.37	4.72	4.67	3.85	4.3	4.84			
35	MAMMA1002371	7.21	3.57	4.06	7.96	12.17	10.93	5.47	3.81	6.44	*	+	
	MAMMA1002380	6.65	2.95	5.07	7.2	8.08	10.65	3.09	4.7	4.45			
	MAMMA1002384	4	1.78	2.02	5.31	7.82	7.61	2.14	4.39	2.73	*	+	
	MAMMA1002385	1.81	2.58	0.88	2.71	5.37	2.61	2.77	1.86	3.22			
	MAMMA1002390	7.22	4.09	4.3	4.23	4.19	5.43	8.27	6.12	7.86			
40	MAMMA1002392	6.65	3.55	1.7	3.98	7.13	4.08	2.98	3.25	3.05			
	MAMMA1002396	10.94	5.98	7.24	14.33	18.89	22.98	6.91	9.41	11.76	*	+	
	MAMMA1002399	6.9	2.88	1.85	8.11	6.41	8.49	4.7	4.28	4.05			
	MAMMA1002400	1.74	0.88	0.89	1.88	3.53	2.38	2.6	2.64	0.96			
	MAMMA1002409	4.98	2.45	2.94	3.65	3.94	4.37	3.81	6.25	5			
45	MAMMA1002411	5.54	2.15	1.5	3.44	5.65	4.97	2.26	3.08	1.74			
	MAMMA1002413	12.21	5.64	2.48	9.88	11.9	8.93	6.13	5.59	4.64			
	MAMMA1002417	3.93	2.05	1.27	4.37	4.53	3.05	1.96	4.22	3.47			
	MAMMA1002427	6.03	2.26	2.41	5.84	9.22	5	5.51	3.52	6.07			
	MAMMA1002428	3.76	1.67	1.82	4.3	5.95	5.66	4.02	3.17	2.93	*	+	
50	MAMMA1002433	8.04	2.9	2.73	4.67	5.92	6.23	3.94	2.95	5.38			
	MAMMA1002434	8.11	3.72	2.87	9.52	10.57	9.58	3.29	5.17	4	*	+	
	MAMMA1002446	3.79	2.83	2.72	3.64	5.3	4.09	2.21	3.52	3.36			
	MAMMA1002447	6.44	2.97	3.54	5.33	7.97	7.41	2.58	4.01	4.02			
	MAMMA1002454	19.95	10.05	7.32	23.49	19.29	16.59	13.96	10.08	15.17			
	MAMMA1002461	12.83	8.73	5.05	6.25	10.29	8.7	5.37	6.47	8.29			
55	MAMMA1002463	8.41	6.54	4.81	4.6	7.71	7.29	6.73	4.8	6.72			
	MAMMA1002464	7.42	5.06	2.53	4.57	5.16	4.31	6.56	4.89	5.9			

Table 222

	MAMMA1002041	2.83	1.69	0.23	3.14	3.37	3.59	1.39	2.49	3.1			
	MAMMA1002042	5.88	3.59	2.24	4.97	5.99	7.54	2.94	3.98	4.72			
5	MAMMA1002045	2.41	1.74	1.47	5.35	8.87	6.75	3	4.53	2.32	**	+	
	MAMMA1002047	5.33	2.17	2.82	3.83	6.17	6.04	1.68	3.55	2.24			
	MAMMA1002056	12.39	6.58	4.37	20.56	18.36	19.17	8.24	9.27	8.66	**	+	
	MAMMA1002058	6.27	2.84	3.39	8	8.2	9.71	5.08	4.13	6.51	*	+	
	MAMMA1002060	1.5	3.41	0.94	1.36	1.83	1.14	1.54	1.23	1.52			
10	MAMMA1002065	9.08	4.91	4.66	8.35	11.05	9.12	3.27	5.48	5.84			
	MAMMA1002068	6.34	2.81	1.47	4.59	6.64	9.1	3.39	3.22	5.73			
	MAMMA1002070	4.29	2.1	1.76	2.92	4.72	3.16	2.15	3.57	3.06			
	MAMMA1002078	5.04	2.14	3.64	3.66	4.1	4.18	2.08	3.2	5.45			
	MAMMA1002080	6.83	3.54	2.1	2.95	4.44	2.95	2.06	5.27	3.19			
15	MAMMA1002082	8.06	4.39	2.39	7.44	9	7.6	3.58	5.19	3.55			
	MAMMA1002084	5.52	4.28	3.59	5.1	6.35	5.81	3.08	4.41	3.89			
	MAMMA1002087	2.38	2.18	1.81	1.76	3.43	2.93	2.59	2.65	3.27			
	MAMMA1002091	5.42	7.29	2.65	4	6.91	4.49	4.2	3.64	5.26			
	MAMMA1002093	1.93	2	0.58	5.96	1.9	2.8	1.65	1.71	2.83			
20	MAMMA1002095	5.4	2.74	3.59	3.25	4.43	4.61	2.69	3.88	4.12			
	MAMMA1002108	5.49	3.13	2.43	2.96	4.71	4.19	2.48	1.84	3.62			
	MAMMA1002112	2.09	1.02	0.93	2.26	2.09	1.19	0.86	2.05	1.87			
	MAMMA1002118	4.48	1.67	0.26	1.23	3.74	1.59	0.63	2.22	1.71			
	MAMMA1002119	8.58	4.34	2.71	5.72	6.62	5.85	3.59	5.08	6.24			
25	MAMMA1002125	9.57	5.01	5.66	13.06	12.09	12.55	6.22	5.68	8.12	*	+	
	MAMMA1002126	13.46	5.9	6.29	18.17	24.01	20.42	8.52	7.83	10.14	*	+	
	MAMMA1002128	5.36	2.96	2.77	3.71	5.08	4.6	3.95	3.22	4.97			
	MAMMA1002132	10.12	4.97	5.63	12.89	10.87	14.39	10.04	6.43	10.71	*	+	
	MAMMA1002140	1.72	1.95	1.35	4.11	5.59	3.44	1.38	1.98	2.23	*	+	
	MAMMA1002142	6.23	4.13	6.33	4.88	8.41	5.57	2.7	5.34	6.44			
30	MAMMA1002143	7.91	3.86	1.2	4	8.63	6.78	4.54	4.01	8.01			
	MAMMA1002145	12.14	5.89	4.12	12.19	9.19	9.27	7.73	5.23	7.12			
	MAMMA1002147	4.21	2.54	2.46	6.44	4.91	6.18	4.06	3.93	4.81	*	+	
	MAMMA1002153	5.55	2.41	3.01	3.35	4.54	5.5	3.13	4.08	5.58			
	MAMMA1002155	9.29	6.93	5.81	15.05	16.47	13.36	7.79	8.57	9.36	**	+	
35	MAMMA1002156	0.5	0.43	0.34	1.18	0.77	0.53	0.87	1.99	2.58			
	MAMMA1002158	3.36	2.26	1.87	4.83	4.63	4.78	2.02	3.6	3.09	**	+	
	MAMMA1002164	4.2	5.9	2.06	5.48	5	6.18	2.35	2.71	6.87			
	MAMMA1002165	9.16	4.19	3.07	5.86	7.65	9.97	4.78	4.68	8.08			
	MAMMA1002170	2.61	1.94	1.29	2.52	2.68	1.48	2.55	4.49	2.09			
40	MAMMA1002174	4.84	4.21	3.36	9.26	11.06	9.43	3.61	5.85	5.69	**	+	
	MAMMA1002175	3.66	3.08	1.47	4.24	3.36	3.13	3.56	5.23	4.15			
	MAMMA1002180	9.95	5.24	8.36	6.25	12	9.82	8.31	11.32	10.45			
	MAMMA1002198	7.77	3.94	4.6	11.59	10.97	8.42	5.79	8.09	5.83	*	+	
	MAMMA1002205	6.94	2.43	4.08	12.68	10.23	10.6	4.99	6.19	5.86	*	+	
45	MAMMA1002206	4.97	3.21	3.83	3.93	5.39	5.02	3.15	4.77	5.02			
	MAMMA1002209	5.93	1.39	2.1	5.8	6.14	5.58	2.65	3.01	4.62			
	MAMMA1002215	25.36	13.93	13.82	17.32	25.36	18.76	19.04	14.22	18.26			
	MAMMA1002219	6.6	5.08	3.39	6.83	8.53	7.54	5.44	5.14	6.2			
	MAMMA1002224	8.1	9.24	5.62	14.79	19.7	17.59	7.17	10.07	8.16	**	+	
50	MAMMA1002229	3.07	2.57	2.61	4.9	4.15	4.71	3.87	4.96	3.11	**	+	
	MAMMA1002230	5.84	5.63	4.35	11.67	10.96	14.46	5.06	7.28	7.47	**	+	
	MAMMA1002233	5.99	1.67	2.56	4.66	5.13	4.71	1.73	5.03	4.75			
	MAMMA1002234	2.42	2.28	2.06	6.51	4.38	3.03	2.11	2.84	3.32			
	MAMMA1002236	9.04	9.45	4.47	5.41	11.26	4.51	4.88	5.38	10.34			
	MAMMA1002243	5.3	1.99	1.09	3.09	2.98	3.83	2.89	2.41	4			
55	MAMMA1002250	6.06	6.45	2.48	6.45	6.62	8.63	6.12	5.22	8.76			
	MAMMA1002253	25.92	17.49	11.68	17.95	18.93	21.68	17.92	18.81	17.12			

Table 221

	MAMMA1001773	6.61	3.09	3.86	5.22	5.33	3.63	5.11	4.68	6.54			
	MAMMA1001778	4.17	2.72	2.42	4.48	7.37	5.12	3.01	4.78	4.22			
5	MAMMA1001783	6.42	4.36	3.89	10.62	14.19	14.31	4.67	8.23	6.82	**	+	
	MAMMA1001785	8.22	2.97	-5.14	14.68	12.34	15.26	7.67	8.51	8.54	**	+	
	MAMMA1001788	2	0.87	0.27	0.81	1.38	1.73	1.53	0.58	0.8			
	MAMMA1001790	5.36	3.86	1.92	6.66	16.36	9.58	3.91	3.27	3.37			
	MAMMA1001800	3.52	2.19	1.41	1.85	4.05	2.73	1.44	1.76	1.56			
10	MAMMA1001804	6.25	3.82	2.87	4.53	3.88	4.64	4.42	4.04	3.96			
	MAMMA1001806	3.43	3.08	1.93	7.24	8.78	6.25	3.11	4.51	5.23	**	+	
	MAMMA1001812	2.22	1.53	1.51	2.28	2.36	2.64	1.38	2.87	1.34			
	MAMMA1001815	1.3	0.41	0.62	2.99	1.2	2.47	2.3	2.24	1.48		*	+
	MAMMA1001817	1.37	3.74	1.14	2.04	2.4	3.09	1.01	1.65	1.29			
15	MAMMA1001818	2.76	5.34	1.53	1.82	5.05	3.5	2.09	2.95	4.34			
	MAMMA1001819	5.52	3.47	3.12	6.33	7.32	6.74	3.51	2.89	5.62	*	+	
	MAMMA1001820	2.45	1.25	0.82	2.09	2.1	3.98	4.93	5.44	3.89		**	+
	MAMMA1001824	6.23	3.21	3.26	6.85	6.39	6.61	3.99	4.27	4.97			
	MAMMA1001832	3.67	1.55	1.58	4.4	5.34	6.5	1.89	2.88	2.54	*	+	
20	MAMMA1001836	7.21	6.9	2.37	8.79	8	7.74	7.22	5.59	4.27			
	MAMMA1001837	8.71	5.61	5.12	7.73	9.45	10.52	4.01	4.19	6.46			
	MAMMA1001848	3.49	1.69	1.44	2.63	4.08	4.52	1.91	2.78	1.99			
	MAMMA1001850	20.05	8.18	11.43	18.79	13.27	17.94	12.58	9.7	17.74			
	MAMMA1001851	6.25	2.81	2.47	7.34	6.62	10.7	4.31	3.59	5.68			
25	MAMMA1001852	7.89	5.2	4.18	14.68	10.33	12.24	6.74	5.9	7.65	*	+	
	MAMMA1001854	8.11	3.75	3.83	5.47	8.12	7.92	4.25	4.74	5.11			
	MAMMA1001858	5.29	6.33	3.33	4.8	9.86	6.77	4.43	4.52	4.66			
	MAMMA1001864	6.57	3.87	3.53	5.26	5.92	6.2	4.84	4.25	4.74			
	MAMMA1001868	7.13	2.35	1.77	6.07	8.46	12.04	4.49	2.72	4.43			
30	MAMMA1001874	2.56	0.8	0.99	1.13	2.27	2.32	0.71	0.85	1.82			
	MAMMA1001878	14.71	6.24	5.55	12.93	17.25	13.98	8.14	7.86	10.4			
	MAMMA1001880	8.73	3.97	3.36	7.33	11.41	9.31	6.98	4.88	7.07			
	MAMMA1001885	8.89	4.03	4.1	9.41	9.07	9.64	3.45	4.7	8.89			
	MAMMA1001890	10.42	4.8	4.27	13.94	12.16	12.45	5.05	4.52	6.53	*	+	
	MAMMA1001893	8.64	3.63	4.1	6.16	5.52	7.2	5.63	4.73	6.76			
35	MAMMA1001901	3.39	1.13	2.13	3.15	3.75	4.39	2.43	2.45	3.16			
	MAMMA1001907	12.12	8.44	5.76	15.43	12.7	15.66	5.86	7.16	6.54	*	+	
	MAMMA1001908	16.6	10.48	11.12	10.97	16.32	14.93	6.4	9.69	8.54			
	MAMMA1001919	1.82	0.17	0.6	0.94	1.34	0.71	1.26	0.88	0.98			
40	MAMMA1001931	3.36	2.44	1.38	2.23	3.72	3.2	2.14	2.05	2.86			
	MAMMA1001937	5.76	3.91	4.17	7.43	4.75	5.56	4.86	3.34	6.3			
	MAMMA1001951	9.42	4.25	4.02	11.76	11.79	12.88	6.81	5.98	6.3	*	+	
	MAMMA1001956	12.62	6.26	4.43	11.46	11.33	13.51	7.86	7.63	5			
	MAMMA1001957	7.69	6.91	2.97	9.44	10.13	11	3.86	6.71	4.82	*	+	
	MAMMA1001960	8.09	4.17	5.2	8.83	7.29	10.11	4.77	4.56	4.66			
45	MAMMA1001963	1.4	0.45	0.94	0.59	0.92	1.24	0.53	1.56	0.54			
	MAMMA1001969	14.58	7.72	8.73	21.99	28.29	25.27	9.69	8.49	9.83	**	+	
	MAMMA1001970	13.52	3.54	5.52	13.53	17.34	15.88	8.28	8.88	8.45			
	MAMMA1001978	1.45	1.06	0.2	0.2	0.85	0.8	1.52	1.12	0.55			
	MAMMA1001992	10.84	5.7	4.65	11.47	10.31	11.31	7.27	6.07	8.17			
50	MAMMA1001994	10	5.97	3.81	5.9	10.24	11.51	7.66	6.84	4.2			
	MAMMA1002000	4.32	3.45	1.54	2.22	2.63	3.21	2.43	4.92	3.14			
	MAMMA1002009	6.14	4.06	3.61	6.87	8.92	11.78	3.75	4.94	3.85	*	+	
	MAMMA1002011	7.71	3.01	4.35	2.88	2.54	4.06	2.17	2.53	2.34			
	MAMMA1002022	5.37	5.17	1.74	4.7	10.07	6.42	3.41	3.09	3.25			
55	MAMMA1002024	16.93	11.72	9.52	17.19	15.15	15.52	13.59	12.95	15.19			
	MAMMA1002032	11.99	7.54	5.7	14.48	16.59	16.42	8.97	9.05	9.69	*	+	
	MAMMA1002033	7.72	10.65	3.5	9.82	12.49	7.85	5.26	4.36	7.58			

Table 220

	MAMMA1001534	1.04	1	0.48	0.51	0.82	0.82	0.58	0.71	1.3				
	MAMMA1001535	4.92	2.88	1.16	1.88	3.67	4.55	1.49	2.38	2.87				
5	MAMMA1001547	6.61	3.6	2.98	6.07	6.82	8.95	4.29	5.11	5.04				
	MAMMA1001551	6.07	3.86	3.57	4.63	5.65	6.3	4.24	3.97	4.09				
	MAMMA1001569	3.5	1.48	2.2	2.86	2.79	2.47	2.33	2.98	1.96				
	MAMMA1001575	8.12	4.85	4.3	5.13	5.29	4	4.97	4.91	5.14				
	MAMMA1001576	20.26	7.19	9.68	8.21	9.38	6.87	9.09	8.98	9.31				
10	MAMMA1001584	4.62	2.36	1.31	4.08	5.15	3.32	1.55	1.67	4.4				
	MAMMA1001586	1.88	3.47	0.76	1.07	3.5	1.99	1.25	2.13	3.7				
	MAMMA1001590	12.7	4.74	4.76	9.14	12.67	13.3	5.6	5.77	7.89				
	MAMMA1001599	4.33	1.21	1.88	2.45	2.99	4.36	2.56	2.76	2.04				
	MAMMA1001600	5.33	1.77	2.89	2.89	5.09	5.36	2.48	3.86	2.92				
15	MAMMA1001604	7.87	5.11	1.45	4.32	5.42	5.07	3.4	3.25	5.07				
	MAMMA1001606	9.46	4.93	4.75	9.09	8.64	10.49	4.91	6.03	6.85				
	MAMMA1001609	2.95	1.2	1.3	2.12	2.38	3.64	2.68	2.56	2.15				
	MAMMA1001614	4.39	2.53	1.88	2.49	3.22	3.59	2.48	3.41	3.61				
	MAMMA1001615	6.67	1.9	1.82	2.35	2.21	3.65	2.11	2.71	3.83				
20	MAMMA1001619	19.31	10.08	12.63	10.87	10.6	14.3	14.55	8.6	14.72				
	MAMMA1001620	8.92	3.44	4.44	6.63	10.03	12.83	4.85	6.18	5.9				
	MAMMA1001623	3.58	4.58	2.08	1.56	2.91	2.34	1.28	2.13	2.52				
	MAMMA1001626	2.57	1.13	1.2	1.48	2.12	1.89	1.75	2.77	3.1				
	MAMMA1001627	2.24	1.39	0.54	2.13	3.22	2.88	2.13	2.52	2.05				
25	MAMMA1001630	3.02	5.98	2.09	4.38	4.01	5.45	2.54	3.3	3.8				
	MAMMA1001633	6.31	4.02	1.66	8.75	9.37	5.34	5.49	3.61	5.08				
	MAMMA1001634	8.31	4.18	4.46	11.22	16.21	13.47	7.21	6.09	6.17	*	+		
	MAMMA1001635	8.83	4.02	2.32	12.04	8.31	8.32	5.06	3.5	2.52				
	MAMMA1001649	4.06	1.62	1.65	3.2	3.67	3.34	1.61	2.68	2.21				
30	MAMMA1001654	7.5	5.7	4.13	5.16	7.53	6.42	3.33	5.51	3.69				
	MAMMA1001660	28.42	20.01	15.26	32.5	33.59	28.79	16.52	14.53	17.32				
	MAMMA1001663	16.19	8.13	7.37	24.06	22.04	19.25	11.83	9.81	14.91	*	+		
	MAMMA1001670	6.04	4.74	3.32	6.72	7.02	6.98	4.35	4.11	5.69	*	+		
	MAMMA1001671	3.01	0.89	1.27	2.72	3.99	2.13	1.77	2.54	1.32				
	MAMMA1001679	4.8	3.29	3	3.03	4.77	2.84	4.71	2.51	4.64				
35	MAMMA1001683	6.21	3.81	4.22	11.62	10.92	14.02	7.47	6.25	5.71	**	+		
	MAMMA1001686	1.2	1.06	0.86	1.34	1.65	3.46	1.07	2.23	3.61				
	MAMMA1001688	27.08	14.53	17.18	23.31	26.84	30.3	37.53	34.87	43.95		*	+	
	MAMMA1001689	10.7	4.3	2.46	5.85	12.72	6.26	3.96	2.83	5.24				
	MAMMA1001692	5.97	3.39	4.03	11.66	13.26	13.23	4.66	4.11	3.69	**	+		
40	MAMMA1001711	7.12	3.2	3.17	7.6	8.99	7.95	4.59	5.62	7.5				
	MAMMA1001715	5.07	1.86	2.28	7.77	5.67	4.34	3.14	3.85	3.95				
	MAMMA1001730	5.56	2.96	1.32	1.82	2.04	2.43	2.03	3.01	2.56				
	MAMMA1001735	17.93	11.2	11.92	16.49	13.17	19.36	14.97	10.91	15.84				
	MAMMA1001740	2.62	1.39	2.19	3.94	5.07	3.81	2.69	2.45	2.08	*	+		
45	MAMMA1001743	63.77	35.5	45.41	34.01	34.01	44	19.91	22.06	23.3		*	-	
	MAMMA1001744	1.18	0.45	0.11	1.34	1.3	0.81	0.46	0.4	0.67				
	MAMMA1001745	12.45	7.1	4.31	14.99	16.74	16.98	8.77	5.37	11.73	*	+		
	MAMMA1001751	5.01	2.42	3.03	4.8	5.52	7.04	3.9	3.22	3.1				
	MAMMA1001752	15.56	8.33	10.02	13.09	14.3	13.11	10.96	9.67	11.14				
50	MAMMA1001754	5.78	4.59	3.53	9.06	6.92	8.14	9.82	5.67	8.59	*	+		
	MAMMA1001757	1.64	0.65	0.62	1.81	1.16	1.05	0.91	2.59	1.38				
	MAMMA1001760	15.19	8.82	7.01	15.51	12.28	21.03	9.85	11.53	17.24				
	MAMMA1001764	2.52	1.27	1.35	2.11	2.1	2.28	1.29	2.52	2.02				
	MAMMA1001767	3.67	2.6	1.45	4.72	4.48	6.08	3.4	1.82	3.79	*	+		
	MAMMA1001768	3.4	1.95	1.15	4.85	4.7	4.24	2.45	3.01	3.52	*	+		
55	MAMMA1001769	10.2	4.54	6.07	16.86	18.38	16.58	8.55	6.27	9.46	**	+		
	MAMMA1001771	7.06	9.36	4.23	3.92	5.03	5.23	5.55	6.69	8.65				

Table 219

	MAMMA1001271	18.48	7.38	8.91	10.48	14.14	10.31	9.58	8.4	12.04				
	MAMMA1001274	4.43	3.8	2.81	4.94	7.96	7.95	4.24	5.07	5.33	*	+		
5	MAMMA1001280	1.75	0.68	1.07	1.62	2.08	1.61	1.59	2.67	1.12				
	MAMMA1001283	7.51	3.83	-5.22	4.97	9.33	8.85	4.6	3.72	6.27				
	MAMMA1001284	9.53	6.17	5.52	8.14	10.72	8.86	5.52	7.53	9.35				
	MAMMA1001286	24.45	16.7	10.97	12.09	12.45	13	5.97	6.39	7.92				
	MAMMA1001289	8.47	4.9	3.19	5.53	5.66	7.55	4.68	4.32	5.33				
10	MAMMA1001292	6.67	3.9	2.86	4.2	6.48	5.22	3.23	4.27	4.41				
	MAMMA1001296	7	4.06	4.91	10.25	16.18	16.77	6.43	5.53	5.34	*	+		
	MAMMA1001298	4.11	3.91	3.07	8.57	9.18	8.84	4.16	3.98	4.76	**	+		
	MAMMA1001305	5.35	2.58	3.48	7.15	5.55	7.22	4.85	4.29	6.18	*	+		
	MAMMA1001309	1.7	1.52	0.97	5.04	3.61	5.38	2.09	2.69	2.33	**	+	*	+
15	MAMMA1001310	10.44	4.9	7.15	8.11	11.15	12.2	4.69	5.11	6.27				
	MAMMA1001322	2.58	0.43	0.4	1.79	2.43	1.77	1.08	1.1	1.59				
	MAMMA1001324	4.35	2.2	1.73	3.98	4.2	3.7	1.97	2.42	3.42				
	MAMMA1001330	13.9	7.33	5.29	11.99	11.29	12.32	6.89	5.13	7.87				
	MAMMA1001333	10.64	5.27	5.22	12.45	17.04	13.72	7.27	8.49	8.86	*	+		
20	MAMMA1001334	19.83	12.61	11.33	16.84	18.12	18.43	11.85	9.39	18.95				
	MAMMA1001337	6.8	2.68	3.43	4.92	5.69	6.15	4.3	5.31	5.13				
	MAMMA1001341	3.94	2.12	2.51	4.82	3.58	4.32	2.93	4.08	4.66				
	MAMMA1001343	4.64	4.02	3.95	10.45	11.27	11.13	3.36	5.55	6.66	**	+		
	MAMMA1001344	3.2	1.52	0.8	2.99	5.13	4.05	4.81	3.84	5.02		*	+	
25	MAMMA1001346	3.61	1.95	1.75	2.88	2.78	3.94	2.71	2.77	4.61				
	MAMMA1001383	13.98	5.18	5.89	17.88	22.89	19.58	10.5	8.81	9.31	*	+		
	MAMMA1001388	6.8	2.8	3.94	7.53	10.07	7.51	5.93	5.82	6.51				
	MAMMA1001396	11.03	6.21	4.6	12.55	13.22	12.6	7.14	6.44	7.15	*	+		
	MAMMA1001397	8.15	4.45	6.77	11.06	10.6	9.93	5.76	7.2	5.97	*	+		
30	MAMMA1001401	12.38	7.29	6.74	14.61	13.5	16.44	10.3	14.7	12.59	*	+		
	MAMMA1001408	3.01	1.06	1.25	3.39	2.85	2.94	2.29	2.63	3.03				
	MAMMA1001411	13.87	6.35	6.18	6.44	8.45	4.19	7.07	7.42	10.12				
	MAMMA1001414	8.9	4.02	3.1	8.97	5.29	6.61	6.05	4.52	6.79				
	MAMMA1001415	10.6	3.71	5.04	5.41	5.06	7.32	4.77	5.68	6.24				
	MAMMA1001418	5.7	2.73	2.09	6.08	5.21	5.62	4.02	2.75	3.87				
35	MAMMA1001419	4.73	2.65	2.23	4.77	8	8.11	4.53	3.83	4.07	*	+		
	MAMMA1001420	3.1	2.15	1.27	3.76	5.4	5.17	2.79	4.4	3.79	*	+		
	MAMMA1001426	18.02	14.05	10.52	23.03	29.5	27.85	14.93	16.81	15.67	*	+		
	MAMMA1001428	19.49	9.42	10.79	23.13	21.75	19.76	15.67	13.18	13.4				
	MAMMA1001432	11.31	4.42	3.74	13.45	13.13	13.68	6.17	5.31	10.64	*	+		
40	MAMMA1001435	5.17	2.46	1.9	6.79	5.64	6.54	4.02	2.35	4.67	*	+		
	MAMMA1001442	5.06	2.93	3.93	6.1	7.84	8.67	6.15	4.58	6.02	*	+		
	MAMMA1001446	12.46	5.86	4.49	8.24	8.89	13.91	4.69	4.66	5.57				
	MAMMA1001450	4.63	2.5	2.67	4.93	4	5.12	3.59	2.97	3.49				
	MAMMA1001452	6.13	3.91	3.22	5.79	9.5	8.17	5.22	5.47	4.79				
45	MAMMA1001465	26.46	18.98	20.83	12.75	32.75	41.93	22.64	25.99	25.3				
	MAMMA1001476	5.04	2.17	1.67	4.15	3.25	3.38	3.37	3.42	3.65				
	MAMMA1001478	8.65	3.83	3.78	10.05	11.02	9.81	4.96	6.35	7.03	*	+		
	MAMMA1001479	12.55	5.38	4.01	10.03	11.12	10.85	9.53	8.55	11.85				
	MAMMA1001487	3.39	1.73	3.53	4.32	4.6	4.59	2.05	2.41	4.9				
50	MAMMA1001498	9.96	8.14	3.99	14.3	13.63	9.71	5.08	12.98	6.54				
	MAMMA1001501	10.61	5.97	4.92	6.54	6.18	6.58	4.88	5.96	6.07				
	MAMMA1001502	8.18	4.06	3.9	5.74	5.38	7.37	5.92	4.78	5.08				
	MAMMA1001510	2.96	0.75	0.46	1.67	1.4	1.25	0.55	1.62	1.38				
	MAMMA1001522	5.03	2.4	1.29	4.2	3.19	3.32	3.17	2.13	2.87				
	MAMMA1001529	6.71	2.99	3.35	4.53	4.35	5.16	2.95	3.56	4.27				
55	MAMMA1001532	9.52	5.54	4.9	8.06	8.4	11.77	6.31	5.77	5.73				
	MAMMA1001533	5.96	3.56	2.76	3.85	3.07	5.41	3.42	3.85	4.91				

Table 218

	MAMMA1001037	9.85	4.28	3.71	10.53	13.73	9.2	7.98	5.87	7.42				
	MAMMA1001038	3.03	1.45	2.07	4.49	7.26	6.95	4.49	3.88	6.41	*	+	*	+
5	MAMMA1001041	6.12	4.31	3.78	4.26	5.32	5.37	5.37	4.53	3.75				
	MAMMA1001043	9.46	4.63	3.66	5.68	7.75	7.15	4.92	3.72	4.88				
	MAMMA1001050	6.35	5.89	3.9	5.29	10.15	10.16	5.02	6.56	5.49				
	MAMMA1001054	5.51	4.13	3	8.5	8.45	8.15	5.21	3.63	4.46	**	+		
	MAMMA1001059	15.39	8.08	6.23	9.1	11.74	11.86	8.44	7.77	9.49				
10	MAMMA1001066	16.43	8.7	9.35	16.38	15.95	15.31	10.1	8.21	12.62				
	MAMMA1001067	3.67	2.44	1.56	5.04	5.4	5.91	3.35	3.05	4.31	*	+		
	MAMMA1001072	11.88	5.32	6.63	6.72	4.61	6.46	5.54	4.86	5.86				
	MAMMA1001073	5.21	2.94	1.75	2.04	3.72	2.45	1.94	3.03	2.39				
	MAMMA1001074	3.99	4.38	2.27	4.13	9.96	13.79	3.27	3.81	5.24				
15	MAMMA1001075	5.54	2.96	3.2	3.06	7.9	7.5	2.62	3.18	3.18				
	MAMMA1001078	7.94	4.65	4.05	9.11	13.65	11.41	7.34	5.68	7.64	*	+		
	MAMMA1001080	22.36	9.18	10.44	11.87	12.56	12.61	9.96	10.5	13.83				
	MAMMA1001082	4.52	3.3	1.66	3.03	5.82	3.36	3.3	2.6	2.45				
	MAMMA1001091	0.73	0.99	0.34	1.07	1.55	1.04	1.3	1.37	1.5		*	+	
20	MAMMA1001092	3.38	1.71	1.14	4.68	5.06	3.84	2.72	2.57	3.2	*	+		
	MAMMA1001094	23.07	10.75	8.74	19.47	15.51	11.95	11.1	12.09	9.06				
	MAMMA1001105	8.97	7.82	3.9	7.84	13.25	10.97	5.27	6.89	7.2				
	MAMMA1001110	1.34	0.28	1.07	0.83	1.4	1.91	0.64	1.83	0.87				
	MAMMA1001126	11.76	5.19	6.22	18.27	20.42	20.62	10.8	7.93	10.63	**	+		
25	MAMMA1001133	13.96	7.98	6.29	17.52	21.82	18.6	12.41	9.09	11.57	*	+		
	MAMMA1001139	16	10.86	8	75.48	52.51	90.41	4.72	2.94	4.09	**	+	*	-
	MAMMA1001141	3.54	2.73	2.73	3.35	3.24	4.02	3.37	4.28	4.25				
	MAMMA1001143	9.1	5.11	2.81	6.09	8.1	8.79	3.94	3.97	7.09				
	MAMMA1001145	8.33	4.95	3.62	3.46	6.81	6.75	3.46	5.11	7.05				
30	MAMMA1001150	8.4	3.25	2.79	2.57	3.1	4.61	3.41	4.01	4.33				
	MAMMA1001154	10.09	4.99	5.59	11.85	11.71	18.3	6.93	7.19	6.3				
	MAMMA1001159	9.34	6.32	4.92	5.06	4.86	4.07	3.31	2.7	4.01				
	MAMMA1001161	14.59	7.23	8.28	17.47	24.12	19.35	11.34	7.11	8.84	*	+		
	MAMMA1001162	8.3	3.74	4.22	6.24	6.6	5.21	4.88	5.43	5.84				
35	MAMMA1001181	5.83	2.22	1.87	4.38	4.79	3.53	3.65	3.3	3.3				
	MAMMA1001186	7.43	2.73	2.8	9.55	11.46	10.04	5.94	5.12	6.23	*	+		
	MAMMA1001189	5.2	2.45	3.28	2.21	6.23	8.54	2.7	3.48	4.97				
	MAMMA1001191	7.35	3.89	3.31	3.72	5.24	6.78	3.27	4.86	5.76				
	MAMMA1001198	420.1	187.9	245.8	305.4	416.1	499.3	169.9	159.8	188.3				
40	MAMMA1001202	22.54	12.72	10.05	25.35	28.4	25.81	14.74	13.68	16.11	*	+		
	MAMMA1001203	10.49	4.64	4.15	9.25	14.44	10.45	6.11	7.56	8.28				
	MAMMA1001206	4.15	2.67	2.33	5.52	7.44	5.57	3.53	2.86	3.88	*	+		
	MAMMA1001208	6.57	2.81	3.7	5.42	5.59	5.39	4.2	3.8	4.35				
	MAMMA1001215	10.79	5.58	5.27	10.75	14.22	15.01	5.67	7.42	7.48				
45	MAMMA1001220	9.93	5.68	4.3	14.65	18.62	17.06	7.53	7.5	9.1	**	+		
	MAMMA1001222	1.59	0.92	0.2	0.96	1.98	1.96	-0.04	1.13	0.9				
	MAMMA1001223	4.89	1.72	1.83	2.87	4.51	4.18	2.3	4.01	2.37				
	MAMMA1001232	8.78	2.9	3.18	7.54	10.45	9.18	4.93	4.96	6.51				
	MAMMA1001234	7.4	4.59	2.41	6.32	6.84	8.88	3.78	3.73	6				
	MAMMA1001237	2.49	1.76	1.72	1.22	2.15	2.66	1.99	1.8	2.56				
50	MAMMA1001243	2.36	1.9	1.62	4.41	7.15	5.33	4.22	3.46	4.95	*	+	**	+
	MAMMA1001244	2.4	1.42	0.68	1.53	3.03	2.44	2.05	2.71	3.6				
	MAMMA1001249	5.06	0.96	1.74	3.77	9.25	4.93	2.44	3.57	4.08				
	MAMMA1001256	2.41	7.77	2.44	2.13	6.99	6.49	2.44	2.38	3.83				
	MAMMA1001259	5.56	2.92	3.02	4.36	6.71	5.33	2.23	3.39	5.03				
55	MAMMA1001260	13.79	6.11	6.31	13.52	13.26	12.23	7.61	6.68	10.66				
	MAMMA1001262	9.64	6.71	5.97	8.72	6.36	5.96	5.41	6.02	8.33				
	MAMMA1001268	4.72	2.75	3.01	9.39	6.71	7.53	3.74	4.95	5.14	*	+		

Table 217

	MAMMA1000855	1.7	1.63	1.03	1.59	2.99	3.96	1.06	2.13	1.04				
	MAMMA1000856	6.3	3.91	3.68	6.66	6.53	6.39	5.69	5.47	5.67				
5	MAMMA1000859	30.54	14.5	21.77	15.43	16.32	21.44	10.77	8.93	11.82				
	MAMMA1000862	3.63	1.84	2.53	2.21	2.9	4.05	1.42	1.82	1.19				
	MAMMA1000863	6.2	3.01	3.04	4.59	9.69	8.1	4.1	6.66	5.5				
	MAMMA1000865	0.8	0.11	0.15	0.67	1.37	0.92	0.2	1.71	0.5				
	MAMMA1000867	4.15	2.15	1.95	2.19	5.49	3.51	1.75	2.5	2.37				
10	MAMMA1000875	9.92	4.24	6.11	6.91	11.92	12.78	4.67	4.48	7				
	MAMMA1000876	4.63	2.26	3.14	3.33	5.28	6.68	4.51	3.48	5.36				
	MAMMA1000877	9.58	4.24	6.31	9.18	13.08	15.47	7.32	6.45	8.51				
	MAMMA1000878	8.16	4.46	5.1	7.91	13.1	10.3	5.72	5.68	6.98				
	MAMMA1000880	4.25	2.2	2.38	4.84	4.93	5.5	2.27	3.49	2.89 *	+			
15	MAMMA1000881	4.86	3.39	4.01	5.58	9.07	9.97	3.7	4.59	4.69 *	+			
	MAMMA1000883	4.1	2.09	3.9	3.29	3.78	3.16	2.41	3.12	3.57				
	MAMMA1000897	0.87	0.78	1.52	1.35	2.84	1.6	1.61	1.81	0.9				
	MAMMA1000898	14.3	5.37	5.9	6.61	8.53	8.2	8.24	7.58	9.2				
	MAMMA1000905	6.32	4.16	3.03	7.58	8.06	10.95	4.06	4.04	6.22 *	+			
20	MAMMA1000906	4.24	2.45	3	4.3	3.89	5.72	2.87	4.2	3.18				
	MAMMA1000908	1.27	0.39	0.86	1.42	2.93	1.74	2.49	2.77	1.87			*	+
	MAMMA1000911	0.41	1.25	0.84	1.86	2.28	2.63	8.08	5.76	7.77 *	+	**	+	
	MAMMA1000914	5.03	2.41	2.68	4.67	4.17	3.32	1.99	2.14	2.33				
	MAMMA1000920	3.12	1.17	2.51	3.63	3.17	3.45	2.05	3.06	3.19				
25	MAMMA1000921	3.37	3.29	3.26	3.61	9.57	6.95	3.48	3.25	3.54				
	MAMMA1000931	8.02	4.92	5.62	10.56	14.6	15.07	6.35	6.66	5.94 *	+			
	MAMMA1000940	6.43	3.57	4.1	8.17	7.42	11.2	5.43	7.24	5.59 *	+			
	MAMMA1000941	8.08	4.42	5.26	11.96	15.08	14.97	7.8	6.29	7.57 **	+			
	MAMMA1000942	16.28	7.28	9.32	16.51	16.66	17.99	9.16	10.49	11.15				
30	MAMMA1000943	8.02	5.62	7.75	12.59	16.34	17.28	9.76	9.93	7.72 **	+			
	MAMMA1000952	8.49	4.92	6.82	13.66	13.4	12.11	7.68	8.43	10.02 **	+			
	MAMMA1000956	1.29	1.15	1.49	1.35	3.18	2.29	2.16	3.08	2.19		*	+	
	MAMMA1000957	6.37	3.36	2.47	7.39	11.27	10.47	4.72	6	5.03 *	+			
	MAMMA1000962	14.04	6.88	6.94	17.04	23.21	26.2	11.63	8.86	12.79 *	+			
35	MAMMA1000966	7.34	3.73	4.5	10.84	15.74	12.34	4.66	6.62	6.12 *	+			
	MAMMA1000968	7.71	3.48	2.83	8.85	11.98	9.01	6.3	7.27	5.97 *	+			
	MAMMA1000972	1.58	1.55	1.15	4.38	2.9	3.02	2.22	4.51	2.3 *	+			
	MAMMA1000973	3.5	1.69	1.59	3.69	3.21	4.33	2.55	2.9	1.2				
	MAMMA1000975	2.22	2.8	2.6	2.48	6.62	3.03	2.24	4.33	2.06				
	MAMMA1000976	7.5	4.17	5.75	10.05	14.48	15.04	6.28	7.31	7.44 *	+			
40	MAMMA1000979	6.1	3.13	2.84	6.83	11.15	7.34	4.03	3.36	5.99				
	MAMMA1000986	8.92	4.73	5.33	9.12	17.71	11.66	6.36	10.27	8.03				
	MAMMA1000987	4.61	3.28	2.96	7.53	9.04	9.57	3.67	3.25	4.14 **	+			
	MAMMA1000988	6.9	4.02	3.13	9.98	9.41	10.85	6.42	4.87	6.36 *	+			
	MAMMA1000994	3.37	2.44	3.14	3.15	4.33	4.9	3.61	4.21	3.97		*	+	
45	MAMMA1000998	3.52	2.26	2.81	4.12	6.42	7.42	3.48	4.56	3.6 *	+			
	MAMMA1001003	1.84	1.4	1.47	5.67	6.98	6.89	2.14	3.71	2.23 **	+			
	MAMMA1001007	0.12	0.01	0.3	0.22	0.03	0.58	0.25	0.21	0.73				
	MAMMA1001008	6.4	6.37	4.3	6.99	5.97	6.01	5.02	5.89	7.81				
	MAMMA1001013	6.8	3.38	4.83	15.25	11.23	8.98	5.96	5.39	9.13 *	+			
50	MAMMA1001014	7.76	3.67	2.44	4.42	6.29	6.7	2.43	2.82	2.35				
	MAMMA1001021	7.09	2.52	2.8	7.68	6.46	6.9	4.64	3.79	3.74				
	MAMMA1001024	8.72	3.44	3.61	8.02	10.11	9.19	4.3	6.16	5.88				
	MAMMA1001025	1.98	1.65	0.42	0.75	1.1	1.07	0.62	0.65	0.73				
	MAMMA1001028	3.61	3.77	2.41	1.41	2.09	2.3	1.65	2.01	1.32		*	-	
	MAMMA1001030	3.45	1.67	2.14	3.47	2.37	4.44	2.07	2.57	2.47				
55	MAMMA1001035	13.14	8.77	7.89	19	23.71	18.79	11.21	8.37	14.57 **	+			
	MAMMA1001036	11.51	6.94	5.48	11.14	14.27	13.18	7.47	5.06	7.52				

Table 216

	MAMMA1000652	8.47	3.81	5.01	8.32	13.85	13.05	5.34	6.27	6.14				
	MAMMA1000657	5.07	3.94	3.85	6.77	10.59	9.73	5.63	6.14	5.11	*	+		
5	MAMMA1000664	2.69	1.1	1.96	4.49	4.5	4.2	2.58	4.29	2.35	**	+		
	MAMMA1000667	4.79	1.98	2.15	4.21	4.93	5.76	3.08	4.06	3.71				
	MAMMA1000668	2.4	1.13	1.67	3.73	2.97	3.09	0.95	4.13	2.08	*	+		
	MAMMA1000669	1.17	0.4	0.79	2.08	2.59	2.37	1.24	0.92	0.96	**	+		
	MAMMA1000670	7.56	4.44	3.7	4.32	4.44	6.75	2.59	5.1	5.48				
10	MAMMA1000672	7.79	2.99	3.4	4.22	3.53	5.63	3.72	4.19	6.43				
	MAMMA1000681	4.68	1.14	3.03	2.41	2.85	4.06	2.7	2.22	3.58				
	MAMMA1000684	35.85	22.61	24.91	21.42	31.5	29.68	12.4	13.65	15.36		*	-	
	MAMMA1000696	6.4	3.52	4.51	7.83	11.25	15.25	8.55	6.27	7.54	*	+		
	MAMMA1000702	8.51	4.05	5.46	6.26	5.22	7.23	5.02	5.02	4.55				
15	MAMMA1000706	3.68	1.19	1.86	2.9	2.36	3.42	2.81	1.88	2.14				
	MAMMA1000707	3.62	1.77	1.28	1.62	3.45	1.98	2.41	2.52	2.5				
	MAMMA1000713	5.4	2.54	3.24	5.36	5.73	6.33	4.52	4.76	4.87				
	MAMMA1000714	7.46	4.12	5.15	8.57	7.81	8.68	8.73	7.85	8.07				
	MAMMA1000718	3.29	2.59	1.62	6.31	6.72	5.21	3.55	3.17	4.84	**	+		
20	MAMMA1000720	11.1	3.49	5.25	10.45	13.49	12.85	6.43	5.97	7.74				
	MAMMA1000723	2.28	1.69	2.12	4.14	3.59	4.23	2.79	2.97	1.93	**	+		
	MAMMA1000731	1.86	0.62	0.69	2.69	3.19	3.37	2.54	2.31	2.78	*	+	*	+
	MAMMA1000732	4.46	2.1	1.55	3.27	6.08	6	3.73	4.07	3.22				
	MAMMA1000733	2	0.47	0.64	1.76	2.5	2.33	0.99	1.71	0.41				
25	MAMMA1000734	19.84	13.3	8.71	14.98	15.8	18.61	13.99	14.24	10.98				
	MAMMA1000736	12.43	4.93	6.22	7.65	6.62	9.44	6.16	4.05	8.82				
	MAMMA1000738	9.86	3.76	4.66	5.29	7.95	8.71	4.04	5.76	4.24				
	MAMMA1000744	6.53	4.63	4.71	11	10.23	11.31	6.26	6.39	7.29	**	+		
	MAMMA1000746	1.48	2.11	1.07	4.85	6.59	5.04	2.55	4.44	6.76	**	+		
	MAMMA1000748	9.39	7.13	8.61	8.38	10.56	16.11	5.63	9.36	9.45				
30	MAMMA1000751	19.32	15.21	15.9	12.13	17.33	24.65	8.32	12.47	10.06		*	-	
	MAMMA1000752	4.99	3.06	2.62	6.31	5.93	7.52	3.57	3.3	3.21	*	+		
	MAMMA1000757	16.42	7.46	8.63	15.03	20.13	20.42	10.82	9.38	12.45				
	MAMMA1000760	13.83	4.85	6.07	16.93	20.12	21.36	9.26	10.09	9.12	*	+		
	MAMMA1000761	7	5.05	5.28	10.4	11.63	13.03	5.86	6.75	6.32	**	+		
35	MAMMA1000775	4.08	1.66	2.88	3.15	4.48	7.4	3.92	4.45	3.2				
	MAMMA1000776	6.7	4.59	3.36	9.35	9.08	9.79	6.68	5.65	5.84	*	+		
	MAMMA1000778	5.98	3.45	2.59	7.46	6.58	10.39	4.17	4.75	3.98				
	MAMMA1000781	5.48	3.83	3.81	4.84	4.93	5.96	2.78	5.06	3.06				
	MAMMA1000782	15.43	7.59	9.38	7	8.75	12.93	6.89	10.66	10.04				
40	MAMMA1000784	6.69	3.02	3.41	4.23	8.26	6.49	8.78	3.6	3.47				
	MAMMA1000788	18.64	7.23	10.16	10.95	9.2	11.24	9.78	6.25	8.61				
	MAMMA1000798	2.84	1.31	1.28	2.57	6.45	2.47	2.42	2.49	2.05				
	MAMMA1000802	10.19	4.79	5.55	11.64	14.85	12.54	8.45	6.23	7.37	*	+		
	MAMMA1000810	10.4	4.83	5.83	11.45	14.19	14.79	8.3	8.84	9.48	*	+		
45	MAMMA1000813	3.06	1.41	1.3	0.97	1.08	1.47	1.17	2.87	1.61				
	MAMMA1000814	11.43	4.36	6.48	10.9	13.12	14.78	6.64	8.56	8.44				
	MAMMA1000824	4.94	1.4	2.5	6.51	8.16	10.38	6.57	7.55	6.92	*	+	*	+
	MAMMA1000827	5.81	3.08	3.37	6.5	5.83	6.58	3.91	3.77	4.74				
	MAMMA1000831	3.49	2.19	2.43	2.04	2.83	2.54	2.49	2.54	3.51				
50	MAMMA1000838	7.72	7.34	6.75	10.55	7.02	15.37	8.46	6.62	9.4				
	MAMMA1000839	9.86	5.11	5.3	13.32	14.94	15.98	11.39	9.61	11.68	**	+		
	MAMMA1000841	2.16	2.22	2.46	2.34	3.62	2.61	2.07	3.51	3.1				
	MAMMA1000842	9.7	5.15	5.18	5.26	8.56	8.54	4.59	6.92	6.8				
	MAMMA1000843	1.45	0.52	0.63	1.44	1.42	1.66	1.24	1.97	1.12				
55	MAMMA1000845	2.99	0.85	1.74	2.17	3.73	3.02	1.45	3.21	2.14				
	MAMMA1000851	12.84	5.8	5.26	10.17	14.4	13.52	7.61	8.15	8.58				
	MAMMA1000854	5.64	2.1	2.3	6.34	4.33	5.81	6.81	5.87	6.68				

Table 215

	MAMMA1000421	7.88	5.58	3.16	7.31	11.57	11.97	5.34	5.28	5.81				
	MAMMA1000422	4.93	2.9	1.84	2.34	3.07	4.44	2.35	2.99	4.14				
5	MAMMA1000423	3.67	2.88	1.35	2.17	3.71	4.12	2.5	2.73	2.24				
	MAMMA1000424	0.47	0.75	0.45	1.27	1.37	1.76	1.14	1.64	1.04	**	+	*	+
	MAMMA1000429	32.94	14.89	22.85	23.37	29.25	34.16	20.87	25.24	25.81				
	MAMMA1000431	7.98	3.3	4.81	7.45	10.34	14.08	4.8	6.35	5.12				
	MAMMA1000432	4.6	2.09	3.06	2.28	3.64	3.72	3.43	4.61	3.15				
10	MAMMA1000437	6.14	5.61	6.7	6.37	13.88	6.85	9.07	7.74	6.38				
	MAMMA1000444	10.06	5.02	5.92	12.4	21.04	12.87	7.66	9.16	10.21				
	MAMMA1000446	5.86	2.32	2.37	3.48	5.41	5.04	2.92	3.11	3.2				
	MAMMA1000449	5.06	1.88	4.07	4.87	7.02	6.19	3.35	3.99	3.47				
	MAMMA1000457	3.42	1.31	1.57	3.54	3.24	3.66	3.14	3.48	3.29				
15	MAMMA1000458	3.87	1.25	2.08	2.19	3.1	2.93	2.24	2.82	1.85				
	MAMMA1000468	1.49	0.06	0.79	0.79	1.06	1.13	0.34	1.08	0.62				
	MAMMA1000472	11.38	4.74	6.91	9.55	12.61	11.92	6.13	7.53	8.61				
	MAMMA1000473	5.96	3.57	3.53	12.63	7.19	13.81	5.26	5.18	5.28	*	+		
	MAMMA1000477	5.82	2.74	2.51	5.72	8.15	7.58	3.74	4.02	3.75				
20	MAMMA1000478	9	4.17	4.73	12.94	18.52	17.59	8.49	7.88	8.95	*	+		
	MAMMA1000483	14.86	5.67	8.42	11.14	12.83	12.05	7.76	6.25	5.28				
	MAMMA1000490	3.41	1.2	1.17	3.21	2.92	3.1	1.71	2.32	2.64				
	MAMMA1000496	2.46	1.87	1.02	2.44	3.29	2.49	1.44	3.16	1.85				
	MAMMA1000500	1.56	0.84	0.9	2.28	2.75	1.98	1.08	1.9	1.36	*	+		
25	MAMMA1000501	11.66	5.38	5.27	11.85	14.49	13.05	6.88	6.5	10.43				
	MAMMA1000503	1.33	0.54	0.92	1.59	1.74	1.27	1.8	2.54	1.09				
	MAMMA1000506	12.82	9.48	10.39	12.58	12.2	12.4	9.73	8.88	12.24				
	MAMMA1000510	7.01	5.28	6.34	4.55	7.48	6.55	5.31	5.02	4.6				
	MAMMA1000515	7.48	2.78	3.25	5.65	6.45	7.72	3.13	3.76	3.48				
30	MAMMA1000516	5.84	1.9	2.82	5.98	7.85	7.2	2.82	3.57	3.21				
	MAMMA1000522	2.27	1.18	1.41	3.64	3.92	3.54	1.42	3.62	1.52	**	+		
	MAMMA1000524	7.63	2.43	4.92	8.34	11.81	13.33	5.04	5.34	4.54	*	+		
	MAMMA1000528	1.85	0.58	1.07	2.05	2.46	2.53	1.6	1.39	1.82	*	+		
	MAMMA1000534	2.5	1.5	1.3	2.79	2.83	2.9	2.6	2.21	1.6	*	+		
35	MAMMA1000541	10.98	5.23	5.03	6.32	9.31	8.45	6.48	6.33	7.6				
	MAMMA1000550	4.4	3.04	2.74	4.35	5.4	3.92	4.73	3.37	2.94				
	MAMMA1000556	1.48	1.03	1.14	1.83	2.63	2.37	0.93	2.78	1.93	*	+		
	MAMMA1000559	4.37	1.96	1.73	4.8	7.23	5.02	4.99	3.84	3.11				
	MAMMA1000565	4.72	1.49	2.86	6.83	6.65	5.82	4.27	3.68	2.63	*	+		
	MAMMA1000567	3.83	3.37	3.67	5.22	7.17	6.61	3.18	4.82	3.63	*	+		
40	MAMMA1000576	15.99	9.01	6.07	17.4	30.24	29.01	12.9	10.14	12.06	*	+		
	MAMMA1000582	5.54	2.74	3.08	4.19	5.56	6.62	5.53	3.7	3.87				
	MAMMA1000583	4.38	2.28	1.5	5.07	4.75	6.13	4.1	3.32	3.54				
	MAMMA1000585	3.99	1.32	2.85	5.97	7.85	8.52	3.94	4.82	4.06	*	+		
	MAMMA1000587	3.21	2.47	2	4.38	5.07	2.06	5.51	4.86	2.27				
45	MAMMA1000591	3.28	1.11	2.12	2.42	2.51	3.46	1.69	4.06	3.09				
	MAMMA1000594	6.52	3.99	5.77	13.18	11.99	15.24	6.18	7.35	5.58	**	+		
	MAMMA1000597	21.18	8.64	13.27	24.68	32.8	31.71	15.42	19.85	17.87	*	+		
	MAMMA1000605	15	7.83	7.51	16.1	27.17	27.84	15.98	12.24	13.96	*	+		
	MAMMA1000612	7.9	2.22	3.52	4.29	4.53	4.73	3.74	1.84	4.78				
50	MAMMA1000614	21.9	15.16	16.51	11.47	18.81	18.51	16.91	14.24	15.41				
	MAMMA1000616	0.69	0.1	0.08	2.78	1.16	2.29	1.88	2.45	1.31	*	+	*	+
	MAMMA1000621	3.29	2.06	2.49	3.22	4.74	3.92	2.54	4.56	2.58				
	MAMMA1000623	3.66	0.62	3.18	3.6	1.78	3.6	1.68	2.93	1.92				
	MAMMA1000625	21.85	13.69	19.79	20.91	20.47	21.13	14.29	16.77	23.93				
	MAMMA1000635	0.42	0.29	0.29	1.14	0.49	1.07	0.07	1.98	0.45				
55	MAMMA1000643	3.78	2.57	1.76	4.32	6.22	6.75	3.82	3.69	4.44	*	+		
	MAMMA1000646	10.28	5.04	4.34	5.25	6.7	11.98	4.93	10.71	4.89				

Table 214

	MAMMA1000204	7.62	4.53	5.82	9.51	6.8	8.54	5.56	4.69	5.2				
	MAMMA1000207	6.14	2.58	4.15	4.25	4.43	7.49	4.2	4.63	3.58				
5	MAMMA1000214	3.73	2.36	3.5	6.05	6.43	8.36	3.87	5.19	4.02 *	+			
	MAMMA1000220	3.64	2.49	2.27	4.02	3.64	4.91	4.36	4.51	3.83		*	+	
	MAMMA1000221	4.11	1.84	1.12	2.33	12.39	3.34	2.86	3.69	1.65				
	MAMMA1000226	3.4	1.09	2.76	2.96	2.31	2.84	1.92	4.54	2.53				
	MAMMA1000227	5.88	3.58	3.47	4.08	7.55	8.07	3.93	3.56	5.9				
10	MAMMA1000230	6.36	3.63	3.36	3.79	7.14	7.18	4.32	4.39	3.89				
	MAMMA1000241	5.23	2.78	2.92	6.17	10.99	9.16	5.63	6.94	5.92 *	+	*	+	
	MAMMA1000245	71.79	48.41	41.99	49.62	55.47	70.51	36.86	32.29	42.56				
	MAMMA1000248	10.75	5.11	8.19	10.32	13.93	13.73	8.64	7.83	9.87				
	MAMMA1000251	4.47	3.42	3.86	6.07	8.71	10	3.62	6.05	5.41 *	+			
15	MAMMA1000254	2.89	1.15	1.35	4.06	5.79	5.07	1.95	5.72	2.71 *	+			
	MAMMA1000257	7.12	4.26	6.71	11.96	14.47	16.44	5.81	9.74	10.27 **	+			
	MAMMA1000262	12.13	6.11	6.35	9.28	17.3	14.89	11.45	12.94	13.68				
	MAMMA1000264	1.54	1.94	1.06	2.96	5.16	6.26	1.9	2.25	1.92 *	+			
	MAMMA1000266	1.41	0.76	1.44	2.49	3.39	2.45	2.4	2.54	1.43 *	+			
20	MAMMA1000270	8.33	3.85	6.34	9.35	14.72	13.36	5.23	6.67	8.24 *	+			
	MAMMA1000271	3.79	2.55	1.83	6.46	5.81	4.43	3.8	4.01	4.5 *	+			
	MAMMA1000277	2.17	1.07	1.86	2.66	2.04	3.91	1.48	2.33	1.37				
	MAMMA1000278	2.46	1.53	1.53	2.26	1.74	1.78	1.61	3.39	1.57				
	MAMMA1000279	4.53	3.12	3.68	7.71	9.92	13.85	2.86	4.21	4.62 *	+			
25	MAMMA1000283	2.8	0.74	1.34	2.2	3.06	3.24	2.27	2.64	2.53				
	MAMMA1000284	7.09	3.1	3.89	5.31	5.61	7.3	4.33	4.12	6.21				
	MAMMA1000287	3.34	1.37	2.39	5.26	5.17	6.99	4.97	3.06	4.33 *	+			
	MAMMA1000294	18.13	8.47	8.55	15.55	11.48	16.82	12.33	10.64	11.59				
	MAMMA1000298	1.54	0.71	0.82	0.74	1.91	1.79	1.37	1.29	1.02				
30	MAMMA1000302	5.12	2.71	2.69	5.15	5.37	6.89	4.36	4.77	2.99				
	MAMMA1000303	4	2.05	1.59	2.54	3.44	3.95	1.95	2.67	2.43				
	MAMMA1000305	1.38	0.71	0.71	1.7	2.67	3.22	1.16	1.69	1.13 *	+			
	MAMMA1000307	12.76	5.57	7.52	10.78	17.15	13.46	11.84	12.09	11.6				
	MAMMA1000309	0.76	0.89	1.4	1.06	1.34	1.72	1.77	0.93	1.2				
	MAMMA1000312	1.8	1.04	0.87	1.28	0.56	1.1	1.25	1.47	0.9				
35	MAMMA1000313	2.67	3.77	1.89	3.1	6.23	5.66	3.12	2.28	2.98				
	MAMMA1000331	4.12	2.28	1.93	3.93	3.97	5.29	3.56	3.45	3.82				
	MAMMA1000335	6.16	2.7	3.37	3.54	3.79	3.88	3.68	2.45	3.73				
	MAMMA1000339	3.25	1.33	2.61	3.01	4.9	3.33	2.91	2.77	1.92				
	MAMMA1000340	2.6	1.63	1.41	3.96	4.43	4.29	1.81	3.28	2.22 **	+			
40	MAMMA1000348	3.33	1.48	2.34	6.45	6.9	6.21	5.1	3.51	6.66 **	+			
	MAMMA1000356	8.13	2.7	3.74	9.76	8.55	10.65	5.97	5.34	5.67				
	MAMMA1000358	4.37	2.17	1.44	5.1	4.35	4.38	3.5	3.09	3.71				
	MAMMA1000360	7.72	3.05	2.69	11.41	9.78	10.42	6.57	4.42	6.39 *	+			
	MAMMA1000361	7.91	2.97	4.89	10.45	10.37	13.01	6.44	5.43	7.13 *	+			
45	MAMMA1000363	5.44	2.67	2.71	3.44	2.89	4.74	2.99	2.83	3.16				
	MAMMA1000370	8.4	6.64	6.2	6.19	7.25	6.56	6.68	7.49	4.91				
	MAMMA1000371	6.81	4.41	6.08	4.39	3.58	5.6	4.96	6.77	5.24				
	MAMMA1000372	11.86	4.03	5.98	15.22	16.38	16.77	7.36	6	7.47 *	+			
	MAMMA1000385	4.62	2.3	2.77	5.18	7.04	8.05	4.85	4.48	5 *	+			
	MAMMA1000388	6.44	2.83	3.67	5.65	4.46	4.85	4.91	3.34	5.06				
50	MAMMA1000395	5.17	2.17	2.95	3.65	4.16	4.78	3.21	2.41	3.84				
	MAMMA1000402	7.68	3.41	2.88	9.51	10.11	10.62	5.46	6.68	5.96 *	+			
	MAMMA1000403	6.72	2.73	3.78	6.04	7.7	8.56	4.71	5.83	4.03				
	MAMMA1000410	4.02	2.21	1.56	4.09	5.7	5.12	2.32	3.4	1.98				
55	MAMMA1000413	1.97	0.9	1.1	2.1	2.16	1.61	0.81	1.47	1.21				
	MAMMA1000414	3.35	1.71	2.96	4.73	3.34	2.27	4.52	4.04	1.89				
	MAMMA1000416	14.38	8.87	8.86	11.04	19.59	15.46	12.54	10.99	12.32				

Table 213

	LIVER1000101	3.81	2.12	1.66	2.9	3.56	2.76	4.13	3.57	3.85				
	LIVER1000106	3.32	1.56	1.67	3.52	2.18	3.06	2.2	1.66	2.75				
5	LIVER1000108	2.84	1.54	1.24	2.99	3.68	3.4	2.48	3.48	3.39 *	+			
	LIVER1000115	2.61	1.46	1.12	3.02	3.28	3.44	1.96	2.92	2.86 *	+			
	LIVER1000120	5.02	2.94	2.41	3.82	3	3.25	3.35	2.12	2.66				
	LIVER1000138	4.91	0.99	2.36	1.52	2.93	3.2	2.89	4.4	2.68				
	LIVER1000146	11.83	5.09	5.8	8.13	11.73	11.21	7.01	6.1	7.3				
10	LIVER1000148	11.43	4.5	7.19	7.38	7.37	7.45	6.46	5.27	6.13				
	LIVER1000157	33.53	16.69	18.55	25.58	33.97	31.92	16.84	15.36	18.47				
	LIVER1000161	7.22	4.95	3.61	5.26	5.68	6.24	4.45	3.94	6.08				
	LIVER1000167	4.56	2.13	2.81	3.19	3.07	3.13	1.51	2.38	2.42				
	LIVER1000174	3.84	1.31	1.5	1.69	2.19	2.47	1.08	2.69	2.65				
15	LIVER1000185	6.12	3.35	4.22	3.51	3.56	3.98	2.75	3.21	2.98				
	LIVER1000187	3.26	1.56	0.93	1	1.39	1.74	0.82	3.36	0.61				
	LIVER1000190	1.95	1.11	1.59	1.96	1.59	2.03	2.89	1.41	1.66				
	LIVER1000192	10.65	6.24	5.2	5.75	5.77	6.49	6.06	5.8	6.02				
	MAMMA1000009	5.3	2.68	2.46	6.62	5.77	8.83	4.6	3.23	5.23 *	+			
20	MAMMA1000015	5.84	1.77	1.87	1.64	2.88	3.59	3.38	2.5	2.77				
	MAMMA1000019	5.66	2.6	2.84	4.89	9.82	8.95	3.81	3.64	4.85				
	MAMMA1000020	3.8	3.44	4.09	3.56	8.72	8.06	4.37	4.09	4.16				
	MAMMA1000024	2.87	0.82	0.95	1.1	1.88	2.53	1.55	2.13	2.01				
	MAMMA1000025	4.87	2.19	2.6	4.8	5.71	6.47	3.27	3.34	4.13				
25	MAMMA1000043	10.51	5.09	5.02	14.31	20.26	13.23	7.72	9.62	9.43 *	+			
	MAMMA1000045	1.69	0.97	1.62	2.91	3.36	3.57	3.47	1.81	1.55 **	+			
	MAMMA1000046	6.47	2.08	3.57	6.03	7.6	8.45	5.17	3.75	4.66				
	MAMMA1000055	6	3.15	3.53	2.8	3.48	4.97	5.81	4.07	2.35				
	MAMMA1000057	12.48	5.52	7.03	12.15	20.3	15.59	7.03	7.1	8.26				
30	MAMMA1000060	14.43	7.18	9.91	16.29	13.21	18.23	10.59	9.1	11.91				
	MAMMA1000069	7.73	3.61	4.66	6.69	8.82	10.74	4.08	5.81	4.8				
	MAMMA1000084	9.73	3.57	5.05	11.91	14.34	16.88	5.45	7.65	6.73 *	+			
	MAMMA1000085	3.47	1.96	1.87	2.74	2.35	3.06	1.99	2.32	2.6				
	MAMMA1000092	5.41	2.13	2.26	4.85	6.6	6.02	2.97	4.24	4.71				
35	MAMMA1000096	3.78	3.03	1.78	3.72	4.8	6.47	4.17	3.9	6.06				
	MAMMA1000097	4.13	2.95	3.91	5.52	4.24	6.86	3.6	3.62	3.89				
	MAMMA1000102	5.12	2.21	2.7	5.22	5.81	5.02	2.56	4.65	3.65				
	MAMMA1000103	3.31	1.56	2.28	4.58	6.05	6.54	2.94	4.29	3.37 *	+			
	MAMMA1000106	2.7	1.79	2.13	3.04	5.09	5.41	1.36	3.69	2.27 *	+			
40	MAMMA1000117	2.72	1.52	1.22	1.31	2.51	2.71	0.5	1.62	1.27				
	MAMMA1000118	8.14	2.71	2.77	3.78	7.64	6.37	5.72	5.22	4.29				
	MAMMA1000129	4.52	1.62	2.67	3.35	3.9	5.18	1.94	2.89	2.82				
	MAMMA1000133	4.27	1.92	2.22	2.89	3.17	3.71	2.86	2.72	3.28				
	MAMMA1000134	3.24	1.82	3.24	6.48	6.88	8.35	3.29	3.76	4.59 **	+			
	MAMMA1000139	3.29	2.4	1.31	3.92	4.25	4.14	3.22	2.8	2.68 *	+			
45	MAMMA1000141	3.46	1.27	2.24	4.07	4.79	6.79	1.97	2.52	1.91 *	+			
	MAMMA1000143	2.16	0.91	1.71	2.99	2.74	3.39	1.31	2.55	1.46 *	+			
	MAMMA1000150	10.88	7.04	8	8.79	14.06	12.33	3.84	10.55	5.74				
	MAMMA1000155	10.85	5.54	5.47	9.19	13.85	13.81	7.6	7.75	9.58				
	MAMMA1000163	5.58	3.38	2.67	5.07	6.46	5	2.15	2.84	6.5				
50	MAMMA1000171	7.29	4.5	4.08	8.95	12.01	14.93	6.64	7.02	7.82 *	+			
	MAMMA1000173	6.86	4.32	5.72	5.71	7.66	7.6	5.97	5.63	5.95				
	MAMMA1000175	4.12	1.18	0.23	1.4	1.36	1.53	1.51	3.3	1.81				
	MAMMA1000183	7	6.5	5.17	8.02	15.13	12.18	4.61	6.16	5.98				
	MAMMA1000191	6.82	3.67	4.83	4.54	6	5.86	3.61	5.7	4.78				
55	MAMMA1000192	13.21	7.3	7.84	8.79	11.31	9.83	5.1	9.07	11.26				
	MAMMA1000193	6.03	2.64	1.36	3.73	3.78	4.43	3.35	3.38	4.25				
	MAMMA1000198	11.19	3.7	4.24	11.67	15.53	12.34	7.74	5.71	7.75				

Table 212

	HEMBB1002596	11.09	4.04	5.16	6.59	10.3	10.29	7.09	7.57	6.28				
	HEMBB1002600	3.89	1.64	1.46	3.06	2.9	2.86	2.13	3.4	3.88				
5	HEMBB1002601	4.5	1.39	1.18	5.04	4.66	4.04	3.46	3.02	3.12				
	HEMBB1002603	4.45	2.06	2.73	4.75	4.46	7.27	5.07	4.62	4.32				
	HEMBB1002607	3.19	2.05	1.88	4.13	5.39	8.14	1.36	2.63	2.56	*	+		
	HEMBB1002610	1.6	0.63	1.12	0.91	2.52	2.41	0.43	2.1	1.33				
	HEMBB1002613	5.8	3.36	3.29	5.19	8.86	7.2	3.09	3.91	3.13				
10	HEMBB1002614	1.91	1.05	1.32	2.97	5.34	5.46	7.36	8.1	8.6	*	+	**	+
	HEMBB1002615	6.52	2.3	1.68	3.51	2.94	3.31	2.18	2.84	3.84				
	HEMBB1002617	2.28	1.5	2.31	5.27	5.83	5.57	3.83	2.69	3.29	**	+	*	+
	HEMBB1002623	5.51	3.51	3.7	8.51	8.93	10.54	4.79	3.06	5.59	**	+		
	HEMBB1002624	8.23	4.59	5.1	6.42	9.16	10.04	4.11	4.54	4.4				
15	HEMBB1002631	1.08	1.05	0.85	1.12	1.79	1.91	1.08	2.01	0.67				
	HEMBB1002635	2.64	1.42	1.61	2.73	3.71	3.6	1.53	2.71	1.37	*	+		
	HEMBB1002644	8.49	6.36	7.31	6.79	8.07	10.17	5.35	5.79	6.57				
	HEMBB1002654	5.54	2.29	1.98	4.78	6.75	4.59	5.18	4.74	4.09				
	HEMBB1002661	7.71	3.01	2.12	14.08	5.44	5.88	4.41	4.24	3.58				
20	HEMBB1002663	6.55	2.14	3.41	6.43	8.16	7.85	4.77	5.41	5.8				
	HEMBB1002664	6.6	3.98	5.84	6.11	8.43	8.44	6.92	5.8	5.93				
	HEMBB1002677	0.49	0.35	0.24	0.79	1.17	0.86	0.54	1.89	0.92	*	+		
	HEMBB1002683	4.48	3.9	3.87	8.9	10.99	11.79	5.35	4.88	6.92	**	+		
	HEMBB1002684	1.16	0.65	1	2.27	2.67	2.14	1.24	1.93	1.3	**	+		
25	HEMBB1002686	2.67	1.11	1.21	1.17	1.78	1.98	0.85	2.28	1.79				
	HEMBB1002692	1.09	0.83	0.68	1.18	2.26	3.02	1.37	1.16	1.64		*	+	
	HEMBB1002693	15.96	10.15	10.49	21.46	23.57	25.74	17.35	13.97	17.93	**	+		
	HEMBB1002697	2.36	2.43	3.54	11.69	11.93	8.98	4.98	6.73	4.87	**	+	*	+
	HEMBB1002699	13.26	6.7	7.9	16.74	17.15	20.25	11.78	11.33	10.9	*	+		
30	HEMBB1002702	1.17	1.29	1.36	2.27	1.04	3.55	1.45	4.46	2.44				
	HEMBB1002705	6.1	3.71	4.11	7.64	8.16	7.66	4.07	5.33	4.38	*	+		
	HEMBB1002712	1.15	0.19	1.21	2.36	1.07	1.65	1.32	2.34	0.92				
	IMR321000028	14.59	7.8	9.64	7.27	7.89	8.64	3.38	5.26	3.94		*	-	
	IMR321000031	3.67	1.78	1.78	4.24	3.4	4.34	3.69	3.39	3.59				
35	IMR321000034	24.92	15.48	15.01	18.47	24.81	26.67	19.77	14.09	22.91				
	IMR321000039	17.93	8.99	10.18	11.47	11.22	20.12	13.91	11.79	14.04				
	IMR321000044	0.32	0.19	0.19	0.47	1.02	1.05	0.71	2.7	0.69	*	+		
	IMR321000063	54.36	30.23	33.89	54.62	56.68	67.83	34.49	32.64	37.87				
	IMR321000085	21.71	12.85	13.46	11.07	12.01	16.43	14.38	12.89	14.05				
	IMR321000089	3.32	1.43	2.9	5.84	3.39	4.37	2.16	3.41	3.89				
40	IMR321000091	5.29	4.33	6.45	10.44	10.54	14.12	6.4	9.24	7.99	**	+		
	LIVER1000004	3.29	1.11	1.67	1.51	1.5	1.97	2.55	2.25	2.71				
	LIVER1000008	3.19	0.85	0.9	1.97	1.35	1.87	1.63	1.58	2.33				
	LIVER1000011	7.48	3.96	4.16	3.89	4.34	5.74	4.62	4.33	4.73				
	LIVER1000022	18.53	8.45	9.73	12.74	12.74	14.58	13.3	11.15	13.79				
45	LIVER1000025	7.77	2.12	4.44	3.72	7.23	8.2	3.81	4.34	4.79				
	LIVER1000030	4.56	1.88	1.59	2.3	3.48	3.86	1.46	2.61	2.79				
	LIVER1000045	2.68	1.73	3.56	1.99	4.14	2.47	1.85	3.55	1.86				
	LIVER1000046	6.12	3.21	3.54	3.3	3.9	5.04	5.21	3.87	9.2				
	LIVER1000072	2.92	1.19	0.82	1.98	3.04	1.6	2.51	2.14	2.54				
	LIVER1000077	4.63	3.26	3.43	3.77	4.63	3.6	5.23	4.42	4.9				
50	LIVER1000080	2	1.34	1.23	2.91	3.37	3.99	2.78	3.35	3.42	**	+	**	+
	LIVER1000086	4.56	1.24	1.67	1.64	5.31	2.33	4.25	3.53	3.07				
	LIVER1000092	2.68	1.43	1.4	3.38	2.77	3.26	2.88	3.19	2.12	*	+		
	LIVER1000095	4.08	1.45	1.83	2.66	3.55	3.63	2.08	3.97	1.97				
	LIVER1000097	2.68	0.88	1.06	2.99	2.32	2.56	2.6	1.48	1.37				
55	LIVER1000098	2.82	0.74	1.66	1.13	2.25	2.13	2.82	1.76	2.99				
	LIVER1000100	8.61	3.08	3.61	4.27	5.23	7.01	4.06	5.3	6.22				

Table 211

	HEMBB1002296	19.39	12.59	10.1	13.85	9.77	11.58	17.38	19.09	20.6				
	HEMBB1002300	5.98	2.27	2.27	4.97	4.83	5.06	3.39	2.79	3.87				
5	HEMBB1002302	4.79	2.37	2.24	3.34	4.96	4.22	3.13	3.11	2.5				
	HEMBB1002306	2.53	0.59	1.19	2.95	4.01	3.53	2.16	2.15	1.9	*	+		
	HEMBB1002316	1.37	0.21	1.01	1.05	1.85	1.65	1.5	1.08	0.63				
	HEMBB1002326	9.34	4.41	4.08	6.83	11.8	13.52	5.14	6.95	5.58				
10	HEMBB1002327	3.74	1.52	2.2	3.25	6.69	8.05	1.41	2.57	2.14				
	HEMBB1002329	6.65	2.85	3.03	3.55	3.52	4.81	3.39	4	4.24				
	HEMBB1002340	2.45	1.14	0.8	2.72	7.22	1.38	1.47	2.32	1.56				
	HEMBB1002342	18.78	10.67	11.1	11.48	10.39	11.81	11.37	10.6	12.37				
	HEMBB1002358	8.06	4.65	5.88	8.32	11.43	13.39	7.12	5.37	8.32				
	HEMBB1002359	4.65	2.7	3.21	2.57	3.59	5.52	2.05	3.08	3.75				
15	HEMBB1002364	3.68	2.01	1.94	4.35	5.19	5.12	3.24	2.77	3.18	*	+		
	HEMBB1002366	26.64	15.48	15.83	13.61	16.98	21.16	15.49	15.91	17.68				
	HEMBB1002371	2.23	1.84	1.61	9.83	11.88	12.5	6.86	8.63	8.95	**	+	**	+
	HEMBB1002381	6.41	3.55	2.93	4.03	6.29	6.16	5.19	4.39	5.77				
	HEMBB1002383	10.2	4.93	4.09	9.89	9.52	10.26	9.31	9.32	10.54				
20	HEMBB1002387	11.72	4.82	7.2	7.69	8.97	9.71	6.05	7.95	7.6				
	HEMBB1002409	4.35	2.96	2.55	5.95	6.17	9.26	3.8	3.76	3.86	*	+		
	HEMBB1002413	10.96	4.94	5.84	12.47	15.22	15.46	7.04	7.35	7.5	*	+		
	HEMBB1002415	2.9	1.63	1.04	2.46	1.99	2.7	2.07	2.58	1.35				
	HEMBB1002424	2.41	2.37	3.44	2.94	2.65	5.7	0.8	2.25	2.17				
25	HEMBB1002425	6.05	3.85	3.42	8.18	9.21	12.24	4.22	6.67	5.02	*	+		
	HEMBB1002427	8.18	4.1	4.67	3.14	4.27	5.26	6.03	4.48	3.96				
	HEMBB1002442	12.17	4.35	6.23	11.86	16.23	14.17	10.19	3.68	8.32				
	HEMBB1002447	8.82	3.51	5.23	10.28	11.65	12.71	5.54	5.46	6.69	*	+		
	HEMBB1002453	10.1	3.7	4.44	12.2	12.96	16.06	5.85	7.3	7.02	*	+		
30	HEMBB1002457	8.34	2.86	3.7	8.87	9.3	9.53	4.63	5.01	4.51				
	HEMBB1002458	1.84	0.2	0.83	2.21	1.65	2.32	1.18	4.23	1.59				
	HEMBB1002463	13.99	7.17	7.29	17.97	18.05	22.29	8.48	10.09	10.66	*	+		
	HEMBB1002465	3.55	1.09	2.46	1.87	2.68	3.41	1.36	3	1.53				
	HEMBB1002477	3.8	1.74	1.62	2.44	2.7	2.39	2.93	1.14	1.8				
35	HEMBB1002479	1.35	1.53	2.03	10.77	11.28	12.82	19.91	17.51	11.35	**	+	**	+
	HEMBB1002489	8.63	4.67	4.63	7.48	7.18	7.8	5.28	6.57	5.43				
	HEMBB1002492	2.72	1.93	0.73	4.55	5.38	4.56	3.26	3.14	4.65	**	+		
	HEMBB1002495	5.34	4.27	3.39	5.35	7.91	6.17	5.79	5.24	4.34				
	HEMBB1002502	0.83	0.8	0.28	1.27	3.14	4.39	2.38	2.95	1.77	*	+		
40	HEMBB1002509	0.76	0.61	0.36	0.32	0.93	0.91	0.52	1.26	0.72				
	HEMBB1002510	2.29	0.9	0.49	1.25	0.69	0.67	0.59	1.16	0.95				
	HEMBB1002520	10.96	4.42	7.37	13.08	19.28	16.87	8.43	9.05	9.26	*	+		
	HEMBB1002522	2.46	1.73	4.71	2.71	2.15	2.36	2.66	2.31	4.74				
	HEMBB1002527	9.87	7.21	7.79	8.36	11.1	10.55	7.47	6.16	5.86				
	HEMBB1002530	7.03	2.68	3.29	3.79	4.83	3.48	4.44	3.46	4.55				
45	HEMBB1002531	2.36	2.37	1.2	1.94	1.74	2.82	1.39	2.3	1.35				
	HEMBB1002534	4.63	2.48	3.25	4.66	8.41	8.39	2.99	3.62	3.89				
	HEMBB1002536	2.96	1.03	1.7	1.05	3.49	2.9	1.99	2.14	1.93				
	HEMBB1002544	3.87	12.89	3.66	4.05	4.44	5.77	1.79	5.33	2.36				
	HEMBB1002545	6.5	3.17	3.97	5.87	8.72	7.62	5.47	5.22	6.78				
50	HEMBB1002550	3.53	1.59	2.38	1.73	2.12	4.1	3.45	2.4	2.04				
	HEMBB1002556	8.37	2.84	4.27	10.84	11.6	10.64	5.58	6.3	7.41	*	+		
	HEMBB1002571	11.52	7.77	9.15	11.56	13.65	12.93	12.05	12.33	11.31				
	HEMBB1002579	9.78	5.85	5.85	7.97	13.11	12.32	6.51	5.4	6.55				
	HEMBB1002582	7.48	3.22	3.33	10.72	9.33	10.11	3.01	4.39	4.41	*	+		
	HEMBB1002584	5.81	3.4	4.16	3.75	2.97	2.76	1.46	2.06	1.93		*	-	
55	HEMBB1002587	12.23	4.61	5.2	12.45	17.92	18.78	8.13	9.27	7.5	*	+		
	HEMBB1002590	5.23	2.47	3.26	5.42	7.78	6.92	3.77	4.99	5.06	*	+		

Table 210

	HEMBB1002002	0.83	0.59	1.4	1.42	1.71	2.28	1.59	0.62	1.07				
	HEMBB1002005	8.43	2.74	4.65	11.77	11.48	12.25	6.14	5.42	7.41	*	+		
5	HEMBB1002009	0.77	2.18	1.38	1.25	1.38	2.16	0.85	1.5	0.79				
	HEMBB1002013	2.33	1.35	1.79	1.62	1.81	3.45	1.55	1.11	1.73				
	HEMBB1002015	7.48	4.38	3.67	9.87	8.21	13.87	7.92	7.55	9.97				
	HEMBB1002024	12.18	6.96	6.46	7.22	8.12	8.32	6.32	9.38	7.79				
	HEMBB1002035	3.12	1.84	1.81	4.86	5.45	3.22	1.97	2.8	1.44	*	+		
10	HEMBB1002039	3.05	1.27	3	3.79	6.93	5.61	2.49	3.6	2.96				
	HEMBB1002041	7.09	2.89	3.99	5.42	7.13	7.97	5.81	4.83	6.2				
	HEMBB1002042	7.43	3.78	4.66	7.93	11.47	10.08	5.53	6.71	6.67	*	+		
	HEMBB1002043	4.31	1.3	3	5.84	8.07	8.67	4.27	3.64	4.54	*	+		
	HEMBB1002044	1.54	1.29	1.16	1.41	1.89	1.39	1.48	2.04	1.41				
15	HEMBB1002045	13.56	9.28	9.85	18.7	19.69	19.62	11.33	11.49	14.07	**	+		
	HEMBB1002049	0.94	0.9	1.48	2.03	3.05	3.51	1.86	1.85	1.5	*	+	*	+
	HEMBB1002050	2.63	0.87	2.41	2.24	3.31	3.77	1.82	2.42	2.29				
	HEMBB1002051	2.77	1.42	2.72	3.76	4.08	3.57	1	2.97	1.66	*	+		
	HEMBB1002068	11.05	4.29	3.65	7.71	6.55	7.57	7.7	4.29	6.63				
20	HEMBB1002069	13.1	6.94	8.01	16.77	20.06	18.1	11.13	9.92	13.2	*	+		
	HEMBB1002075	2.31	1.12	2.72	4.01	5.39	4.96	2.61	2.52	2.47	*	+		
	HEMBB1002079	3.29	1.28	2.08	2.22	2.42	2.23	2.53	2.39	1.66				
	HEMBB1002080	1.83	2.55	0.96	2.15	2.98	4.39	1.68	2.81	2.3				
	HEMBB1002082	2.22	1.44	1.38	1.35	2.4	2.6	1.2	1.53	2.07				
25	HEMBB1002084	1.85	1.72	1.75	2.73	3.83	5.21	2.72	3.71	3.91	*	+	*	+
	HEMBB1002088	11.64	8.26	10.3	14.66	19.71	16.32	16.11	15.05	19.56	*	+	*	+
	HEMBB1002092	8.42	4.12	3.19	8.1	10.6	9.29	6.67	5.28	5.88				
	HEMBB1002094	8.51	6.18	7.26	14.48	15.44	15.77	7.48	6.89	8.09	**	+		
	HEMBB1002103	13.1	13.5	12.83	61.49	68.55	57.48	66.63	34.04	51.6	**	+	*	+
	HEMBB1002109	6.77	3.65	4.41	10.27	12.78	11.5	7.97	4.24	7.06	**	+		
30	HEMBB1002115	44.63	28.15	32.39	41.8	53.57	63.47	24.84	22.28	27.42				
	HEMBB1002120	2.22	0.77	1.3	3.55	2.83	2.5	1.74	2.54	1.48	*	+		
	HEMBB1002121	1.32	0.72	1.59	2.14	1.84	1.52	1.15	1.56	1.25				
	HEMBB1002134	29.98	14.03	18.39	22.56	28.18	29.08	20.1	20.18	26.29				
35	HEMBB1002136	5.67	2.48	3.78	3.62	3.43	4.97	3.89	4.13	4.88				
	HEMBB1002138	3.55	2.31	2.47	7.41	6.73	5.61	7.6	5.28	8.06	**	+	*	+
	HEMBB1002139	3.56	2.49	3.1	6.05	5.07	6.19	3.34	5.1	3.14	**	+		
	HEMBB1002141	5.57	2.73	5.33	5.02	6.05	7.64	4.99	5.45	6.15				
	HEMBB1002142	4.26	2.17	2.9	5.21	4.83	7.21	3.06	3.4	2.29				
	HEMBB1002145	2.66	1.68	2.79	4.87	2.84	2.91	1.83	3.33	2.18				
40	HEMBB1002152	2.89	1.29	3.31	6.08	5.5	7.8	2.66	3.88	3.38	*	+		
	HEMBB1002162	4.47	2.09	2.74	4.63	5.63	4.42	2.84	4.52	4.28				
	HEMBB1002173	2.01	1.5	1.53	4.12	5.2	7.12	2.21	2.47	3.85	*	+		
	HEMBB1002189	5.63	4.01	3.4	9.38	12.87	12.35	5.18	5.01	5.41	**	+		
	HEMBB1002190	4.01	6.72	3.24	8.35	6.45	9.57	5.06	3.62	5.05				
45	HEMBB1002193	4.3	2.37	3.54	3.79	4.24	4.57	3.11	3.84	2.85				
	HEMBB1002217	8.31	4.18	4.51	10.88	11.96	11.17	4.63	6.35	5.39	*	+		
	HEMBB1002218	21.17	7.63	13.71	19.12	24.55	22.92	14.78	13.7	19.7				
	HEMBB1002228	4.29	2.39	3.53	7.69	9.04	7.22	3.92	7.05	4.8	**	+		
	HEMBB1002232	2.54	0.96	2.12	5.44	4.77	4.87	2	4.33	3.46	**	+		
50	HEMBB1002245	2.24	0.69	1.25	1.7	1.97	1.97	1.7	1.11	1.6				
	HEMBB1002247	2.78	1.52	2.56	1.84	2.86	2.26	3.27	2.52	2.93				
	HEMBB1002249	8.45	3.73	4.77	12.48	12.32	13.64	6.18	5.35	5.42	**	+		
	HEMBB1002254	2.12	1.02	1.52	4.72	4.67	7.07	3.96	3.27	2.9	**	+	*	+
	HEMBB1002255	0.31	0.16	1.07	0.59	0.84	2.46	0.5	0.92	0.27				
55	HEMBB1002266	1.03	0.51	0.66	4.13	2.54	2.5	1.25	1.72	1.36	*	+	*	+
	HEMBB1002271	56.56	35.65	38.07	20.53	29.91	28.83	14.93	16.36	14.09		*	-	
	HEMBB1002280	1.89	0.47	1.28	2.71	3.38	2.75	1.12	1.95	1.11	*	+		

Table 209

	HEMBB1001788	5.11	2.85	2.49	8.04	8.23	9.77	5.27	5.14	5.83	**	+		
	HEMBB1001793	13.59	3.52	4.92	5.61	7.12	5.14	6.71	5.28	5.84				
5	HEMBB1001797	0.88	0.62	1.95	0.94	0.65	0.97	1.07	1.81	1.9				
	HEMBB1001802	6.5	3.72	4.86	7.5	8.03	6.58	5.93	6.91	6.06				
	HEMBB1001812	5.74	3.61	5.29	9.39	12.73	12.64	5.58	6.99	9.37	**	+		
	HEMBB1001815	20.05	9	15.52	27.98	23.86	26.02	37.42	29.06	44.83	*	+	*	+
	HEMBB1001816	5.07	2.26	3.92	9.09	8.62	9.45	5.29	4.77	4.9	**	+		
10	HEMBB1001831	1.2	0.45	0.53	1.8	1.74	1.99	0.55	2.73	1.28	*	+		
	HEMBB1001834	19.83	12.47	10.64	12.5	19.26	19.83	14.74	13.9	15.71				
	HEMBB1001836	4.06	3.15	2.68	7.01	7.21	7.9	3.1	3.18	4.01	**	+		
	HEMBB1001839	1.83	0.36	0.78	1.33	1.05	1.21	1.58	1.39	1.02				
	HEMBB1001841	4.21	3.05	4.61	6.62	7.34	6.85	8.41	7.68	5.57	**	+	*	+
15	HEMBB1001844	4.31	2.59	2.19	5.78	3.8	4.04	2.62	4.06	3				
	HEMBB1001847	11.75	7.16	10.2	21.65	17.41	24.55	7.68	9.92	9.98	**	+		
	HEMBB1001848	2.73	1.25	1.47	4.72	2.91	3.06	15.56	19.7	16.79		**	+	
	HEMBB1001850	7.3	4.6	5.92	9.74	8.83	8.43	10.59	7.86	13.13	*	+		
	HEMBB1001859	6.4	9.16	9.93	12.13	14.98	16.02	18.07	14.33	23.47	**	+	*	+
20	HEMBB1001863	6.66	2.82	3.58	9.9	10.12	11.35	6.68	3.16	7.05	**	+		
	HEMBB1001867	1.21	1.36	0.82	2.34	2.45	3.53	2.08	1.31	1.98	*	+		
	HEMBB1001868	3.28	1.27	0.26	2.34	1.83	1.98	2.3	1.36	2				
	HEMBB1001869	4.99	3.41	2.47	4.55	8.08	7.57	3.34	3.94	4.05				
	HEMBB1001872	3.4	4.06	0.84	4.75	2.37	1.57	2.65	1.38	2.04				
25	HEMBB1001874	2.47	1.57	1.58	3.42	1.79	3.58	3.5	1.76	2.08				
	HEMBB1001875	1.3	0.4	3.1	2.27	2.57	2.84	2.23	0.73	0.98				
	HEMBB1001880	9.6	4.1	4.24	11.57	10.59	10.4	5.78	4.19	6.8				
	HEMBB1001899	2.12	0.58	0.29	1.53	1.49	1.79	2.01	0.55	1.92				
	HEMBB1001903	4.86	1.84	3.46	4.45	3.55	4.47	5.08	3.38	4.99				
30	HEMBB1001905	6.94	3.72	4.24	3.83	3.28	4.45	3.35	1.95	3.04				
	HEMBB1001906	3.51	0.89	1.09	3.56	2.45	3.39	2.27	3.05	2.44				
	HEMBB1001908	1.61	2.17	1.92	5.17	4.2	3.43	1.41	2.29	3.44	*	+		
	HEMBB1001910	2.88	1.38	0.82	4.07	3.93	6.71	2.4	1.88	3.55	*	+		
	HEMBB1001911	6.98	2.87	4.02	9.07	10.54	12.95	3.98	4.78	7.22	*	+		
35	HEMBB1001915	4.25	1.76	1.83	6.42	5.24	7.19	5.74	2.92	4.49	*	+		
	HEMBB1001921	5.38	3.56	4.5	10.21	11.3	11	5.97	4.64	6.62	**	+		
	HEMBB1001922	3.83	1.35	3.8	5.95	3.77	3.39	3.48	2.3	3.67				
	HEMBB1001925	3.73	2.29	2.11	4.2	3.69	3.62	2.81	2.27	3.72				
	HEMBB1001930	0.59	0.63	0.42	2.23	1.25	1.36	0.41	1.35	1.01	*	+		
40	HEMBB1001944	3.88	3.55	3.94	5.26	8.37	10.06	2.98	4.95	4.6	*	+		
	HEMBB1001945	5.17	3.58	5.47	3.15	4.34	6.51	3.41	6.48	6.46				
	HEMBB1001947	6.49	1.48	5.58	2.11	3.59	4.92	2.72	2.7	2.62				
	HEMBB1001950	6.47	3.08	4.75	4.98	5.8	5.65	5.08	4.12	4.55				
	HEMBB1001952	4.62	1.75	2.38	5.87	7.63	6.22	3.88	3.07	2.9	*	+		
	HEMBB1001953	3.33	1.23	1.69	3.8	4.29	3.6	2.72	2.28	2.79				
45	HEMBB1001957	3.22	1.56	1.85	3.38	4.52	4.53	3.81	1.96	3.18	*	+		
	HEMBB1001959	7.02	7.17	6.24	7.94	4.73	8.54	5.15	5.79	4.06		*	-	
	HEMBB1001962	4.04	1.76	3.14	4.32	4.25	6.26	2	2.46	5.87				
	HEMBB1001967	11.44	5.2	6.57	12.83	15.13	16.73	7.11	8.82	7.6	*	+		
	HEMBB1001973	5.08	2.32	4.1	4.86	6.84	9.36	3.18	5.36	3.55				
50	HEMBB1001978	7.53	3.35	6.01	7.28	6.5	6.97	5.55	5.59	5.54				
	HEMBB1001983	20.88	11.32	14	10.33	15.1	15.82	11.27	9.2	12.69				
	HEMBB1001987	1.67	0.99	0.76	2.21	2.1	2.79	2.2	1.61	1.87	*	+		
	HEMBB1001988	1.86	1.73	2.04	2.83	3.58	3.43	2.01	2.02	1.85	**	+		
	HEMBB1001990	4.65	2.51	4.22	4.26	6.45	6.12	4.77	3.77	5.16				
	HEMBB1001996	2.64	1.19	1.29	1.17	1.43	2.67	1.72	2.23	1.63				
55	HEMBB1001997	4.3	2.22	2.71	5.89	6.32	7.41	2.43	4.16	2.74	*	+		
	HEMBB1001999	15.97	11.41	12.12	8.02	17.07	19.1	5.81	7.78	7.71		*	-	

Table 208

	HEMBB1001548	11.61	4.55	5.07	6.22	6.23	8.02	13.1	5.3	6.57			
	HEMBB1001551	2.02	1.27	1.35	2.89	1.88	2.65	2.33	1.18	2			
5	HEMBB1001555	3.38	2.36	2.27	4.34	5.15	4.75	3.71	2.88	3.52	**	+	
	HEMBB1001562	6.73	3.72	2.72	6.03	4.98	4.88	4.48	4.21	3.29			
	HEMBB1001564	143.7	103.7	84.35	117.1	130.5	149.5	73.73	78.89	81.45			
	HEMBB1001565	4.34	2.01	6.14	5.35	4.71	6.93	2.17	3.81	2.88			
	HEMBB1001569	3.35	1.85	2.92	2.44	1.48	3	2.68	1.66	2.47			
10	HEMBB1001573	4.11	1.78	1.25	2.55	2.8	4.22	1.76	2.85	2.7			
	HEMBB1001585	5.19	3.43	2.13	7.14	9.58	10.48	5.06	4.35	4.95	*	+	
	HEMBB1001586	2.45	1.89	1.57	2.45	2.59	4.79	2.08	2.01	1.8			
	HEMBB1001588	9.91	4.02	1.68	7.84	12.44	10.86	6.94	6.15	5.93			
	HEMBB1001595	2.38	2.13	1.24	3.04	4.7	3.31	4.54	3.91	4.77	*	+	**
15	HEMBB1001596	7.58	3.68	4.12	10.26	11.71	11.73	8.26	5.8	7.17	**	+	
	HEMBB1001599	1.66	1.47	1.01	2.08	1.72	2.54	1.43	2.23	1.83			
	HEMBB1001603	1.5	0.25	0.77	1.78	2.38	2.95	1.47	2.06	1.36	*	+	
	HEMBB1001606	0.98	0.3	0.79	0.72	0.7	0.98	0.73	0.96	0.76			
	HEMBB1001612	7.29	5.01	5.69	10.05	12.84	11.6	6.84	5.75	5.35	**	+	
20	HEMBB1001618	2.21	1.9	1	2.28	2.95	2.82	2.58	3.52	1.79			
	HEMBB1001619	2.74	2.34	1.59	5	7.12	6.26	2.86	3.86	3.26	**	+	
	HEMBB1001623	3.47	2.37	1.26	9.12	1.21	1.26	2.81	2.15	1.28			
	HEMBB1001625	0.39	0.5	0.61	1.56	1.46	2.32	2.13	1.91	2.02	*	+	**
	HEMBB1001630	2.05	0.69	1.57	1.73	2.03	1.92	0.69	0.97	1.11			+
25	HEMBB1001635	2.2	0.75	1.17	3.5	2.23	1.77	1.56	1.05	1.51			
	HEMBB1001637	3.51	1.4	2.57	3.58	4.43	4.86	2.1	2.95	2.6			
	HEMBB1001641	1.95	0.54	0.63	1.54	1.04	1.19	1.35	0.64	1.26			
	HEMBB1001653	5.49	2.4	2.56	5.29	5.68	6.05	3.35	3.68	4.27			
	HEMBB1001665	1.36	1.13	0.8	0.24	0.85	0.87	0.48	0.61	0.56		*	-
30	HEMBB1001666	2.05	1.95	2.11	3.16	2.96	2.94	3.08	3.78	1.71	**	+	
	HEMBB1001667	2.49	2.15	1.55	5.36	1.62	4.96	1.46	1.39	2.66			
	HEMBB1001668	1.24	0.08	2.02	7.77	6.22	7.71	3.16	4.06	4.45	**	+	*
	HEMBB1001669	1.14	0.56	0.64	1.01	1.36	1.96	0.82	0.73	1.12			
	HEMBB1001670	4.9	1.43	3.88	3.76	6.22	5.35	4.26	5.99	6.05			
35	HEMBB1001673	9.43	4.46	3.65	7.18	5.87	10.36	4.73	4.98	5.54			
	HEMBB1001675	4.45	1.52	2.55	2.96	2.17	2.25	2.39	2.98	2.34			
	HEMBB1001679	3.43	1.92	1.36	3.15	2.26	1.5	2.37	3.04	2.3			
	HEMBB1001684	3.34	2.15	1.93	2.33	2.97	3.86	3.7	4.19	3.07			
	HEMBB1001685	0.43	0.79	0.82	2.14	2.22	2.08	1.31	1.84	3.11	**	+	
40	HEMBB1001695	0.91	0	0.49	2.21	2.23	2.38	1.38	2.34	1.74	**	+	*
	HEMBB1001703	8.08	2.9	6.21	6.72	7.83	9.08	5.46	5.54	6.16			
	HEMBB1001704	4.34	1.92	3.68	6.91	10.28	8.29	3.81	4.11	3.01	*	+	
	HEMBB1001706	5.33	4.91	1.92	7.82	8.35	10.07	3.64	3.58	4.65	*	+	
	HEMBB1001707	5.79	3.89	4.11	6.65	6.58	7.83	3.81	4.55	4.31	*	+	
	HEMBB1001717	2.9	1.19	1.54	2.56	2.78	3.16	1.54	2.73	1.87			
45	HEMBB1001731	36.41	33.52	31.32	28.11	26.74	25.14	22.32	14.43	19.67	*	-	**
	HEMBB1001734	3.1	2.92	2.47	4.7	5.73	6.77	4.4	3.15	4.26	**	+	
	HEMBB1001735	2.54	0.66	2.3	5.4	4.73	5.13	2.48	3.07	2.77	**	+	
	HEMBB1001736	5.75	4.06	4.43	6.69	5.77	10.19	4.22	3.36	4.39			
	HEMBB1001747	2.44	0.77	1.23	3.44	4.21	3.48	1.4	1.37	2.22	*	+	
50	HEMBB1001749	8.77	3.39	4.72	11.21	15.68	17.47	7.58	5.43	7.33	*	+	
	HEMBB1001753	7.34	3.22	3.36	7.29	7.53	8.22	6.29	5.11	5.25			
	HEMBB1001756	3.12	1.84	2.45	2.82	2.94	4.26	3.19	1.55	3.2			
	HEMBB1001757	0.84	0.19	0.52	0.79	1.64	1.25	0.88	2.16	1.24			
	HEMBB1001760	1.15	0.71	0.71	1.53	1.28	1.9	0.58	2.3	0.49	*	+	
55	HEMBB1001762	2.92	1.03	2.15	2.66	1.94	2.8	2.3	3.2	2.05			
	HEMBB1001780	11.82	11.49	14.29	11.74	11.14	12.3	6.31	7.75	11.3			
	HEMBB1001785	0.42	0.01	1.19	1.62	1.09	1.43	0.08	1.04	1.6			

Table 207

	HEMBB1001299	11.58	8.15	6.05	9.03	8.73	7.82	6.87	6.29	8.87				
	HEMBB1001302	6.82	4.33	3.28	5.31	5.44	7.1	4.47	4.4	7.2				
5	HEMBB1001304	1.87	0.87	0.83	1.3	1.76	2.94	1.91	2.12	1.37				
	HEMBB1001314	2.52	0.38	1.35	1.89	2.07	2.7	1	1.94	1.9				
	HEMBB1001315	2.2	0.42	0.99	1.6	0.99	1.5	2.82	1.66	1.14				
	HEMBB1001317	5.5	2.93	3.71	6	6.29	9.01	6.12	6.25	7.04		*	+	
	HEMBB1001326	1.44	0.28	0.42	1.14	1.37	0.97	0.93	1.85	0.56				
10	HEMBB1001331	3.49	1.15	3.33	3.16	5.21	4.92	2.94	2.32	3.4				
	HEMBB1001335	2.13	0.58	1.32	2.09	1.33	1.73	1.47	0.84	0.69				
	HEMBB1001337	4.69	2.11	3.26	4.29	6.51	6.35	3.43	4.14	3.13				
	HEMBB1001339	3.42	1.11	1.36	2.82	1.69	2.07	1.52	2.17	1.96				
	HEMBB1001344	2.99	1.77	1.84	2.4	2.3	3.56	2.28	2.27	2.05				
15	HEMBB1001346	3.15	2.58	2.53	3.75	3.57	4.79	2.76	4.39	3.22	*	+		
	HEMBB1001348	1.96	1.25	1.97	4.3	3.56	4.57	1.75	3.51	2.64	**	+		
	HEMBB1001350	2.69	1.8	2.82	11.17	12.83	10.95	7.44	8.11	8.86	**	+	**	+
	HEMBB1001356	1.82	0.34	1.21	2	1.23	1.35	0.99	1.53	1.63				
	HEMBB1001364	1.29	0.93	0.89	1.8	2.27	2.41	2.29	1.24	1.25	**	+		
20	HEMBB1001366	3.41	1.36	1.76	6.29	5.97	7.89	2.97	3.23	3.76	**	+		
	HEMBB1001367	5.44	2.63	4.67	5.82	13.11	9.17	6.34	5.62	5.1				
	HEMBB1001369	1.88	0.36	0.91	2.5	3.44	2.87	2.19	3.7	2.34	*	+		
	HEMBB1001380	3.65	2.5	3.07	8.69	9.13	10.12	4.6	7.63	4.24	**	+		
	HEMBB1001381	7.54	3.35	4.95	9.78	7.21	8.91	5.88	6.12	6.67				
25	HEMBB1001384	2.77	2.23	5.27	4.04	4.7	5.21	2.99	5.46	4				
	HEMBB1001387	1.33	0.72	1.19	2.84	1.92	3.26	0.78	2.08	0.69	*	+		
	HEMBB1001394	2.01	1.22	0.71	4.71	4.19	4.99	2.39	2.44	2.66	**	+	*	+
	HEMBB1001407	3.37	1.49	0.8	2.53	3.21	2.87	4.47	1.2	2				
	HEMBB1001410	1.19	0.14	0.37	0.55	0.79	0.77	0.44	1.14	0.17				
30	HEMBB1001413	2.53	1.15	2.11	4.01	6.2	3.82	2.17	2.18	2.56	*	+		
	HEMBB1001419	3.82	1.67	2	5.53	5.54	4.76	5.16	3.44	3.45	*	+		
	HEMBB1001421	1.55	0.78	1.24	9.94	7.28	9.56	5.74	5.75	4.91	**	+	**	+
	HEMBB1001424	0.54	0	0.28	0.9	0.45	0.6	0.84	1.22	0.47				
	HEMBB1001426	2.45	0.64	1.42	3.9	4.18	3.95	2.09	3.09	1.9	**	+		
35	HEMBB1001429	10.12	5.99	4.62	6.28	4.44	8.1	5.21	7.29	9.1				
	HEMBB1001436	11.8	4.02	6.29	22.88	14.63	21.79	9.57	8.07	10.97	*	+		
	HEMBB1001443	1.46	1.5	1.3	2.55	2.11	3.84	5.74	4.67	5.74		**	+	
	HEMBB1001449	4.24	1.68	1.33	4.21	5.76	5.46	2.38	1.89	2.76				
	HEMBB1001454	4.2	2.22	2.85	4.88	5.14	6.3	1.94	2.02	3.61	*	+		
	HEMBB1001458	4.34	4.36	3.05	7.92	4.69	4.55	3.87	3.06	3.94				
40	HEMBB1001461	2.41	1.63	1.39	3.76	3.78	6.76	3.87	1.93	2.34	*	+		
	HEMBB1001463	4.41	1.84	3.33	6.77	8.03	7.56	3.07	2.66	3.3	**	+		
	HEMBB1001464	1.53	1.48	0.96	1.16	0.81	1	0.81	0.25	1.04				
	HEMBB1001466	1.71	1.2	0.87	3.03	2.72	4.34	2.85	2.09	4.25	*	+		
	HEMBB1001482	3.03	1.42	1.06	1.64	2.18	1.42	2.97	1.16	2.1				
45	HEMBB1001500	2.17	1.05	0.9	2.57	2.02	2.37	1.04	1.45	1.55				
	HEMBB1001505	8.22	5.06	7.49	13.32	13.9	13.27	5.5	6.16	7.01	**	+		
	HEMBB1001521	2.58	1.03	1.95	4.68	3.52	3.79	2.8	2.46	2.3	*	+		
	HEMBB1001527	14.66	7.32	7.32	12.93	16.36	15.19	7.53	11.09	12.62				
	HEMBB1001530	7.24	3.1	6.46	5.19	6.93	5.94	6.69	5.92	5.53				
50	HEMBB1001531	5.66	2.3	2.38	5.05	4.74	5.69	3.58	2.66	2.99				
	HEMBB1001532	2.05	0.38	0.82	1.99	0.87	2.3	1.76	1.25	1.24				
	HEMBB1001535	3.86	2.42	2.26	4.62	4.93	5.74	3.17	2.1	4.36	*	+		
	HEMBB1001536	5.02	2.43	2.77	5.57	4.42	5.08	2.95	2.46	3.39				
	HEMBB1001537	3.43	1.79	1.93	5.9	3.91	6.35	3.35	2.86	3.81	*	+		
	HEMBB1001542	10.24	4.77	6.29	8.68	10.49	11.37	4.75	4.74	4.61				
55	HEMBB1001543	4.42	2	4.45	6.17	7.07	7.41	4.96	3.35	2.51	*	+		
	HEMBB1001547	1.69	0.68	1.1	3.41	2.74	1.36	1.07	2.16	2.08				

Table 206

	HEMBB1001068	7.81	3.48	2.43	5.74	4.82	6.22	5.55	5.34	6.4				
	HEMBB1001082	5.14	1.53	2.93	10.11	5.98	8.43	4.89	3.46	4.79 *	+			
5	HEMBB1001095	14.6	9.13	9.13	9.72	6.9	9.06	5.98	7.72	8.46				
	HEMBB1001096	3.56	1.37	1.54	4.69	5.52	4.24	2.24	1.72	3.53 *	+			
	HEMBB1001101	21.47	17.94	10.93	10.99	11.87	12.38	8.8	9.1	8.37				
	HEMBB1001102	2.77	1.29	0.76	2.93	2.4	3.87	2.39	1.32	2.26				
	HEMBB1001104	5.43	2.94	3.94	9.11	5.73	9.85	5.68	2.83	4.42 *	+			
10	HEMBB1001105	3.73	2.54	3.47	3.95	6.18	9.09	3.39	3.81	3.94				
	HEMBB1001112	8.37	6.64	4.97	5.94	6.55	6.82	6.29	6.97	5.99				
	HEMBB1001113	7.58	3.55	4.62	10.53	11.56	12	7.39	5.47	7.82 **	+			
	HEMBB1001114	7.84	3.54	5.33	11.15	12.39	11.97	6.57	3.79	5.55 **	+			
	HEMBB1001115	12.69	6.52	6.38	8.41	6.32	7.74	8.1	3.98	5.13				
15	HEMBB1001117	1.26	0.59	1.14	3.99	3.99	7.09	4.39	3.19	2.89 *	+	**	+	
	HEMBB1001119	2.73	0.69	1.36	3.27	2.76	3.17	1.69	1.82	2.33				
	HEMBB1001126	17.3	8.41	6.34	12.51	13.52	16.39	9.04	9.96	8.26				
	HEMBB1001133	7.22	2.46	6.43	7.94	11.25	15.48	5.58	5.96	7.46				
	HEMBB1001137	4.69	1.94	2.48	3.07	2.31	3.24	4.3	2.74	3.49				
20	HEMBB1001142	10.97	4.26	5.7	14.69	16.82	16.36	7.91	5.78	10.87 *	+			
	HEMBB1001145	8.34	3.24	4.81	10.74	10.95	12.08	5.82	4.69	6.65 *	+			
	HEMBB1001151	8.95	6.02	5.47	5.12	6.22	5.78	8.53	8.19	8.82				
	HEMBB1001153	5.68	3.55	3.85	6.9	7.36	7.26	5.29	4.07	4.12 *	+			
	HEMBB1001158	5.25	4.46	4.73	8.21	9.2	10.97	4.6	4.37	5.83 **	+			
25	HEMBB1001169	5.93	2.46	2.66	6.12	6.91	7.13	3.71	3.73	4.71				
	HEMBB1001170	2.28	0.23	1.68	2.09	1.33	2.33	1.48	1.17	1.14				
	HEMBB1001175	4.7	2.5	2.14	5.28	3.05	6.25	4.06	3.09	3.56				
	HEMBB1001177	11.32	4.92	7.58	14.33	14.36	15.14	8.51	7.62	8 *	+			
	HEMBB1001182	7.1	3.3	3.03	8.51	7.41	6.84	6.75	4.9	5.74				
30	HEMBB1001192	4.01	1.43	2.59	3.22	2.9	2.65	3.81	3.22	2.43				
	HEMBB1001199	1.24	0.85	1.37	0.51	1.77	3.72	1.58	1.98	1.27				
	HEMBB1001200	0.7	0.28	0.37	0.41	0.29	1.06	0.14	0.69	0.72				
	HEMBB1001208	6.24	1.58	2.41	2.54	3.62	5	2.67	3.31	3.15				
	HEMBB1001209	8.96	2.6	4.27	8.47	9.46	10.64	6.12	3.72	4.78				
35	HEMBB1001210	3.39	3.6	6.25	13.57	15.06	13.24	8.2	7.86	10.28 **	+	*	+	
	HEMBB1001215	56.1	31.37	29.04	36.73	42.52	41.17	25.87	19.36	26.75				
	HEMBB1001217	4.33	2.5	3.14	2.96	3.91	4.21	4.42	3.57	4.01				
	HEMBB1001218	4.39	2.08	2.28	6.07	7.97	8.92	4.93	4.87	4.51 *	+			
	HEMBB1001221	1.61	1.15	0.66	1.21	1.16	1.19	2.11	1.68	0.87				
	HEMBB1001224	2.88	1.37	1.83	3.46	3.87	4.78	1.63	2.85	1.71 *	+			
40	HEMBB1001230	3.6	1.44	3.39	4.28	5.22	5.68	2.22	3.15	2.2 *	+			
	HEMBB1001234	9.13	2.44	8.29	5.98	6.49	5.96	5.83	7.02	6.04				
	HEMBB1001235	5.5	2.57	3.09	3.97	3.82	5.68	4.42	5.01	5.46				
	HEMBB1001237	11.86	5.88	6.73	9.88	9.37	10.19	7.04	5.53	6.3				
	HEMBB1001242	3.75	2.48	2.08	4.97	4.37	4.59	4.13	4.47	3.96 *	+			
45	HEMBB1001244	1.32	1.13	0.4	0.82	0.94	1.53	1.73	1.61	1.2				
	HEMBB1001249	3.12	1.54	0.34	2.25	4.83	2.55	2.05	1.99	2.11				
	HEMBB1001253	6.29	1.42	2.97	13.67	4.84	8.24	2.79	2.84	4.65				
	HEMBB1001254	2.47	0.84	1.05	1.37	2.56	1.79	1.57	2.54	1.52				
	HEMBB1001266	1.23	0.44	1.59	2.72	2.03	1.62	2.12	1.69	5.16				
50	HEMBB1001267	7.87	4.02	4.63	13.5	11.84	13.24	5.84	7.38	7.93 **	+			
	HEMBB1001271	4.61	1.62	1.38	4.06	3.96	5.87	2.53	2.67	2.49				
	HEMBB1001282	6.27	3.11	3.61	3.44	3.72	3.72	3.96	3.68	3.39				
	HEMBB1001287	13.66	7.12	7.62	9.05	8.08	10.38	11.92	6.12	11.75				
	HEMBB1001288	3.65	1.71	2.11	2.38	2.54	2.69	2.09	2.39	1.58				
	HEMBB1001289	10.93	6.03	8.57	15.81	16.55	18.1	8.4	7.7	5.92 **	+			
55	HEMBB1001290	3.6	2.6	2.26	3.2	3.1	5.19	4.49	3.29	2.53				
	HEMBB1001294	2.74	1.82	3.02	1.97	1.99	2.92	2.55	2.49	2.42				

Table 205

	HEMBB1000831	5.58	1.72	2.71	4.5	3.81	4.21	2.23	2.64	2.11				
	HEMBB1000835	4	1.57	1.01	4.73	4.53	5.6	3.04	2.52	2.85	*	+		
5	HEMBB1000840	6.38	3.54	3.15	8.28	10.6	8.97	6.91	4.2	4.08	*	+		
	HEMBB1000848	4.7	2.4	2.04	8.23	8.85	8.6	7.06	5.5	6.33	**	+	*	+
	HEMBB1000852	0.54	0.28	0.27	0.52	0.36	0.24	1.16	0.97	0.61		*	+	
	HEMBB1000857	7.91	6.39	3.23	5.68	6.47	7.09	4.42	3.6	4.37				
	HEMBB1000858	5.33	2.35	2.78	9.3	8.37	8.17	3.94	3.82	2.97	**	+		
10	HEMBB1000867	5.01	2.6	3.3	9.23	10.12	8.69	3.49	5.17	4.45	**	+		
	HEMBB1000870	4.43	1.73	2.81	6.64	6.44	7.5	2.8	3.34	3.99	*	+		
	HEMBB1000876	2.52	1.01	1.78	2.03	2.41	3.32	1.17	1.96	2.6				
	HEMBB1000881	4.52	2.25	2.68	3.85	3.48	4.21	3.8	3.6	3.52				
	HEMBB1000883	1.07	0.87	0.48	2.38	2.52	2.42	1.86	2.24	1.15	**	+		
15	HEMBB1000887	16.17	10.38	8.54	18.39	28.8	26.71	14.31	15.73	15.23	*	+		
	HEMBB1000888	1.52	0.47	0.72	0.71	0.87	1.25	1.08	2.54	2.95				
	HEMBB1000890	4.2	1.91	2.82	6.2	6.22	11.04	3.56	3.57	3.05	*	+		
	HEMBB1000893	3.13	1.95	2.57	3.14	8.44	5.73	3.88	3.35	2.73				
	HEMBB1000900	2.72	1.85	1.78	2.31	2.75	4	1.77	1.83	1.88				
20	HEMBB1000905	7.13	4.79	4.05	6.15	5.33	7.36	6.49	7.74	6.04				
	HEMBB1000908	3.42	1.78	2.53	3.45	3.15	4.99	2.18	3.31	2.95				
	HEMBB1000910	3.27	1.5	0.99	3.5	4.25	4.18	2.64	2.6	2.61	*	+		
	HEMBB1000913	1.53	1.02	1.16	2.35	1.71	3.01	2.43	2.82	3.12		**	+	
	HEMBB1000915	125.5	96.58	90.74	52.7	70.12	78.2	138.4	94.57	151.2	*	-		
25	HEMBB1000917	5.94	3.71	3	10.02	9.8	10.14	6.41	5.43	5.2	**	+		
	HEMBB1000927	3.9	2.3	4.04	2.93	2.18	2.45	3.26	2.61	3.09				
	HEMBB1000932	1.41	0.52	1.78	2.08	2.21	2.86	1.55	1.9	0.46				
	HEMBB1000933	63.34	47.44	31.38	44.11	52.4	49.52	46.54	37.21	45.55				
	HEMBB1000936	7.16	3.79	4.04	4.95	3.87	5.38	3.06	2.19	2.36				
30	HEMBB1000939	9.8	5.4	5.5	8.13	8.11	6.88	7.11	4.16	5.78				
	HEMBB1000941	1.26	1.52	1.91	2.33	1.33	3.43	1.03	2.28	3				
	HEMBB1000947	3.84	2.12	3.17	3.27	3.95	6.16	2.65	3.42	5				
	HEMBB1000954	2.09	0.96	1.77	3.22	2.47	2.01	1.52	2.5	2.09				
	HEMBB1000959	1.47	0.69	1.99	4.15	4.21	5.2	2.08	3.64	2.15	**	+		
35	HEMBB1000973	0.93	0.22	1.08	1.36	1.53	1.02	0.58	1.34	0.88				
	HEMBB1000975	6.35	2.45	2.52	2.87	4.55	4.7	3.97	3.56	3.46				
	HEMBB1000981	1.55	0.65	1.17	2.92	1.74	2.12	1.91	1.15	1.6				
	HEMBB1000985	4.16	2.16	3.38	6.79	6.53	7.43	6.9	5.56	5.46	**	+	*	+
	HEMBB1000991	2.4	0.94	2.24	1.58	2.01	2.39	1.83	3.86	2.04				
	HEMBB1000996	6.16	2.86	5.71	15.05	12.65	14.03	9.39	6.89	7.92	**	+		
40	HEMBB1001000	0.81	0.42	1.96	2.31	1.45	2	2.11	2.4	1.74				
	HEMBB1001004	0.63	0.42	0.74	2.36	1.33	1.9	1.27	2.5	0.58	*	+		
	HEMBB1001008	0.9	0.72	1.22	1.95	1.11	0.92	0.7	1.72	0.82				
	HEMBB1001011	4.86	1.41	1.32	2.52	2.1	3.78	2.71	1.63	2.77				
	HEMBB1001014	5.41	3.41	2.83	4.86	8.33	8.51	5.54	2.65	5.28				
45	HEMBB1001020	3.52	1.22	3.22	5.91	7.22	5.47	4.21	2.46	3.29	*	+		
	HEMBB1001024	3.88	2.55	2.6	4.94	7.97	7.2	4.48	3.54	3.57	*	+		
	HEMBB1001026	4.57	3.08	2.54	5.25	5.33	6.61	2.93	3.4	3.78	*	+		
	HEMBB1001037	2.04	0.83	2.17	4.63	4.48	3.78	3.41	3.94	2.4	**	+		
	HEMBB1001042	2.63	0.37	1.26	3.42	3.22	3.69	2.16	3.39	1.69	*	+		
50	HEMBB1001046	3.55	2.14	2.26	3.89	3.63	3.68	3.15	4.56	3.14				
	HEMBB1001047	5	1.57	1.46	5.39	4.72	4.88	2.39	1.51	4.62				
	HEMBB1001048	8.53	3.68	3.67	9.65	6.39	8.39	5.59	5.14	7.15				
	HEMBB1001051	1.18	0.9	0.65	0.91	1.6	1.29	0.9	1.3	2.48				
	HEMBB1001056	4.02	2.51	1.82	4.56	3.43	4.23	3.26	2.37	3.48				
55	HEMBB1001058	4.62	1.41	2.29	4.81	4.08	5.54	4.01	2.62	3.49				
	HEMBB1001060	1.13	0.14	0.28	1.95	1.91	2.6	0.75	1.53	1.56	*	+		
	HEMBB1001063	4.1	1.41	1.69	3.82	4.69	5.11	3.01	2.79	2.86				

Table 204

	HEMBB1000579	1	0.63	1.23	1.94	1.94	1.61	0.75	2.27	0.83	*	+		
	HEMBB1000585	1.32	0.9	1.33	2.89	2.66	2.35	1.39	2.41	1.82	**	+		
5	HEMBB1000586	5.03	2.33	2.86	4.93	10.49	10.9	3.19	3.33	3.66				
	HEMBB1000589	4.34	3.31	2.32	4.73	9.62	7.86	4.05	4.47	4.07				
	HEMBB1000591	6.2	2.47	3.35	5.53	10.43	9.55	5.26	4.88	5.68				
	HEMBB1000592	3.62	1.12	1.49	3.68	3.48	4.83	5.06	2.83	3.4				
	HEMBB1000593	5.63	3.16	4.14	7.95	8.98	9.6	4.23	4.57	4.71	**	+		
10	HEMBB1000595	9.73	4.88	6.49	11.51	8.83	10.26	5.12	4.65	3.54				
	HEMBB1000598	3.08	2.45	2	3.88	5.18	4.28	2.89	4.3	2.68	*	+		
	HEMBB1000611	1.33	0.64	1.43	2.46	1.17	1.82	0.83	1.24	1.6				
	HEMBB1000617	12.12	5.56	4.61	11.59	16.06	19.06	7.94	6.34	10.85				
	HEMBB1000623	7.8	2.76	2.97	7.01	3.89	6.02	4.57	2.94	6.61				
15	HEMBB1000630	2.59	1.28	1.39	2.17	2.39	2.78	2.69	2.13	3.79				
	HEMBB1000631	10.27	4.76	4.53	6.2	6.77	8.48	8.04	7.46	8.07				
	HEMBB1000632	6.25	2.1	3.02	6.63	6.59	8.13	4.84	4.67	4.51				
	HEMBB1000636	13.35	4.72	8.11	7.29	10	13.28	8.71	9.3	9.58				
	HEMBB1000637	26.51	17.46	16.75	28.37	43.24	52.91	24.53	21.76	22.76				
20	HEMBB1000638	1.76	0.67	1.19	2.95	4.12	4.45	1.31	0.92	1.68	**	+		
	HEMBB1000642	10.59	4.41	5.99	11.15	12.92	13.73	6.73	6.84	9.2	*	+		
	HEMBB1000643	1.65	1.83	1.24	2.38	2.51	3.19	2.28	0.92	1.97	*	+		
	HEMBB1000649	3.91	2.47	2.78	5.9	5.23	6.96	3.56	3.95	5.15	*	+		
	HEMBB1000652	6.02	2.91	2.8	5.46	7.5	7.04	3.21	3.43	4.33				
25	HEMBB1000655	12.28	6.34	8.07	9.28	11.26	11.56	6.56	3.92	6.25				
	HEMBB1000665	1.52	0.76	1.22	2.5	1.48	1.81	2.25	0.85	1.56				
	HEMBB1000668	2.21	0.39	1.35	5.91	7.44	6.43	4.09	4.69	4.22	**	+	**	+
	HEMBB1000671	9.73	3.87	4.11	15	14.71	15.82	8.84	8.17	8.33	**	+		
	HEMBB1000673	2	0.92	2.42	2.06	2.03	2.24	2.77	0.96	1.66				
30	HEMBB1000679	1.96	1.55	2.94	3.03	1.89	3.47	3.49	2.72	4.24				
	HEMBB1000684	10.32	4.72	6.06	13.49	17.19	16.84	8.71	6	9.32	*	+		
	HEMBB1000692	2.42	1.11	1.48	1.94	1.06	1.01	1.68	1.28	1.89				
	HEMBB1000693	6.65	3.11	3.35	5.7	3.46	5.14	5.27	4.98	4.93				
	HEMBB1000705	4.28	2.03	1.45	4.17	5.14	4.6	2.08	2.85	2.66				
35	HEMBB1000706	2.4	0.82	1.33	4.76	1.91	1.69	2.35	1.33	2.24				
	HEMBB1000709	5.9	4.56	2.82	9.88	15.43	11.7	9.92	8.98	12.92	*	+	*	+
	HEMBB1000714	4.07	1.84	2.28	3.51	2.48	3.46	4.34	1.56	2.5				
	HEMBB1000725	3.83	2.12	2.8	3.51	3.57	2.91	4.38	2.17	3.5				
	HEMBB1000726	6.74	3.26	3.37	8.38	10.66	11.11	5.09	6.26	5.9	*	+		
	HEMBB1000729	5.92	3.12	3.67	3.82	5.2	5.28	2.93	3.03	3.74				
40	HEMBB1000738	6.27	2.98	4.84	7.01	7	9.14	5.8	4.68	8.01				
	HEMBB1000749	6.38	4.5	8.03	10.82	12.38	19.82	6.87	7.43	9.13				
	HEMBB1000763	4.28	1.52	4.69	3.87	3.73	4.04	3.58	5.24	3.54				
	HEMBB1000770	2.56	1.54	1.45	4.69	5.02	5.12	3.94	2.82	2.01	**	+		
	HEMBB1000774	4.01	2.16	2.61	6.02	5.76	6.03	4.48	3.56	3.59	**	+		
45	HEMBB1000777	16.82	8.94	10.71	11.64	9.96	10.04	11.16	9.95	10.48				
	HEMBB1000781	4.68	2.51	2.03	4.83	6.62	5.74	2.82	4.66	5.27				
	HEMBB1000788	1.26	1.09	0.22	0.77	1.4	0.96	0.82	1.05	1.38				
	HEMBB1000789	3.3	1.16	1.77	2.42	1.9	2.76	1.89	2.74	1.95				
	HEMBB1000790	4.72	2.05	3.39	5.79	6.37	7.78	3.19	2.91	4.28	*	+		
50	HEMBB1000794	0.97	0.54	1.08	1.04	2.04	2.15	0.72	1.24	1.02				
	HEMBB1000807	7.3	3.23	3.76	7.53	4.81	6.34	3.19	2.77	3.98				
	HEMBB1000809	10.2	3.24	6.13	7.78	12.54	11.13	7.52	8.8	9.69				
	HEMBB1000810	6.83	2.64	2.68	4.19	3.73	4.74	4.18	2.82	5.16				
	HEMBB1000821	3.04	1.01	1.43	1	1.91	2.05	1.27	2.15	1.75				
55	HEMBB1000822	1.16	1.15	0.89	1.14	1.34	1	1.68	1.67	1.68		**	+	
	HEMBB1000826	3.27	2.25	2.9	2.37	8.91	8.1	2.85	5.14	2.76				
	HEMBB1000827	4.04	1.85	2.66	4.07	6.2	5.58	3.55	3.41	2.85				

Table 203

	HEMBB1000307	4.53	1.84	2.11	5.17	5.68	6.34	1.82	4.17	2.46	*	+		
	HEMBB1000309	4.37	1.32	2.88	3.56	4.27	5.98	1.82	3.44	1.73				
5	HEMBB1000312	1.28	2.42	1.55	2.15	2.18	2.23	2	1.79	3.52				
	HEMBB1000317	3.2	2.61	1.78	3.01	2.88	2.59	3.81	2.77	1.93				
	HEMBB1000318	4.73	1.3	2.1	5.96	5.69	5.2	3.19	2.91	3.3				
	HEMBB1000332	1.76	1.25	0.79	0.91	1.05	1.63	1.26	1.46	1.31				
	HEMBB1000335	2.8	1.5	1.13	1.18	3.42	3.3	2.66	1.47	1.27				
10	HEMBB1000336	4.55	1.96	1.92	2.95	2.84	3.92	3.25	2.93	2.41				
	HEMBB1000337	14.36	7.11	10.05	9.07	12	11.79	6.71	8.68	8.74				
	HEMBB1000338	4.54	3.23	3.69	5.82	6.25	7.43	2.29	3.11	3.62	*	+		
	HEMBB1000339	6.86	3.25	2.73	8.08	11.02	9.45	5.52	5.3	4.99	*	+		
	HEMBB1000341	6.67	3.9	3.27	5.51	6.05	5.75	4.88	3.76	5.53				
15	HEMBB1000343	5.14	3.78	3.56	8.73	11.85	8.26	4.26	5.37	4.59	*	+		
	HEMBB1000354	5.87	3.91	3.47	10.81	11.74	10.84	4.26	5.4	6.59	**	+		
	HEMBB1000358	6.98	3.62	4.09	5.18	4.64	6.14	4.86	3.92	4.34				
	HEMBB1000369	3.23	1.7	2.29	3.08	3.51	3.68	1.39	2.56	1.97				
	HEMBB1000373	11.86	5.42	7.78	12.45	14.15	14.43	4.75	5.77	6.52				
20	HEMBB1000374	8.03	4.3	5.09	13.94	16.47	17.13	5.55	9.31	7.38	**	+		
	HEMBB1000376	11.27	4.35	3.91	16.2	18.49	19.55	9.94	8.36	10.29	*	+		
	HEMBB1000383	4.6	2.17	1.96	4.57	3.4	3.45	10.39	7.52	9.9		**	+	
	HEMBB1000391	6.84	4.23	4.83	6	8.02	7.16	4.22	5.21	3.67				
	HEMBB1000399	5.23	1.96	3.15	3.41	3.17	3.69	3.69	3.13	1.81				
25	HEMBB1000402	2.6	1.48	0.94	2.16	3.1	1.88	0.98	2.21	2.08				
	HEMBB1000404	1.75	0.76	1.14	1.48	2.07	2.27	1.05	1.58	1.14				
	HEMBB1000407	1.46	1.26	1.6	1.67	2.46	3.55	0.54	2.33	2.09				
	HEMBB1000420	6.02	3.01	5.42	7.53	9.7	10.11	3.76	5.07	4.73	*	+		
	HEMBB1000430	59.23	34.65	23.06	49.23	46.08	51.49	46.72	34.37	41.23				
30	HEMBB1000434	18.16	8.94	9.74	22.34	23.72	31.12	11.49	11.35	12.88	*	+		
	HEMBB1000438	2.81	0.97	1.46	1.87	3.06	1.59	2.06	2.06	1.78				
	HEMBB1000441	5.61	4.55	3.22	9.46	9.64	11.7	6.15	5.84	7.17	**	+		
	HEMBB1000447	6.8	2.32	3.46	10.82	16.06	18.31	25.43	26.28	30.87	*	+	**	+
	HEMBB1000449	1.31	0.73	0.5	2.05	2.12	2.41	1.36	2.6	1.7	**	+		
	HEMBB1000453	8.09	6.85	8.91	11.38	10.07	15.36	7.99	10.3	12.98				
35	HEMBB1000455	2.98	3.4	2.03	3.63	4.91	3.97	1.67	3.24	1.52				
	HEMBB1000472	7.59	4.06	3.3	4.71	4.91	6.8	5.17	4.42	5.06				
	HEMBB1000480	9.8	3.69	3.57	8.18	11.17	10.77	5.35	5.7	6.17				
	HEMBB1000486	7.07	2.27	3.48	8.16	9.71	10.13	5.36	5.39	6.03	*	+		
	HEMBB1000487	2.41	1.44	1.32	2.02	2.24	3.56	1.77	2.52	2.72				
40	HEMBB1000490	9.25	6.82	8.08	12.41	16.92	19.33	9.89	8.92	10.33	*	+		
	HEMBB1000491	6.31	3.37	4.57	9.52	7.65	10.48	5.02	4.69	3.79	*	+		
	HEMBB1000492	2.22	0.64	1.44	4.93	5.13	7.41	2.99	2.91	3.63	**	+	*	+
	HEMBB1000493	4.06	2.22	4.19	4.24	6.19	6.18	1.66	2.72	2.91				
	HEMBB1000510	6.41	3.47	4.28	6.87	9.13	11.79	5.4	4.78	5.74				
45	HEMBB1000516	4.76	2.42	3.32	9.01	5.12	5.56	4.9	3.1	6.78				
	HEMBB1000518	1.77	0.89	0.96	2.32	1.98	1.84	1.86	2.02	1.38				
	HEMBB1000523	5.6	4.26	4.37	10.14	11.92	12.71	5.32	6.89	8.07	**	+		
	HEMBB1000530	2.95	1.4	1.93	9.88	7.75	9.87	4.5	4.51	2.86	**	+		
	HEMBB1000542	8.28	5.69	6.91	10.8	11.2	12.53	8.2	7.18	7.39	**	+		
50	HEMBB1000550	1.32	0.8	1.53	2.82	2.53	3.26	1.75	3.01	2.05	**	+		
	HEMBB1000554	7.82	3.63	5.16	11.58	14.79	21.33	5.25	6.34	9.2	*	+		
	HEMBB1000556	7.65	3.11	3.74	5.66	6.17	8.22	5.32	4.21	5.38				
	HEMBB1000564	4.88	2.2	2.92	4.79	4.84	5.7	5.46	2.87	3.44				
	HEMBB1000567	11.63	5.99	6.65	15.29	18.3	19.22	9.27	9.36	9.27	*	+		
	HEMBB1000569	5.23	1.99	2.42	5.2	5.06	5.8	7.16	8.5	8.18		*	+	
55	HEMBB1000573	7.84	3.79	4.94	12.01	13.33	13.4	9.04	8.26	7.72	**	+		
	HEMBB1000575	5.33	4.35	4.85	8.19	11.22	12.98	7.01	6.31	6.36	*	+	**	+

Table 202

	HEMBB1000008	6.33	3.99	3.55	9.32	9.9	11.83	4.69	4.68	5.58	**	+		
	HEMBB1000018	9.18	4.31	7.12	14.89	18.9	20.93	7.15	7.95	8.65	**	+		
5	HEMBB1000024	8.61	5.93	3.83	12.18	15.58	14.42	6.22	5.32	8.3	**	+		
	HEMBB1000025	7.18	1.68	2.62	5.76	5.35	5.09	4.63	4.5	5.11				
	HEMBB1000030	5.99	4.74	5.88	11.95	12.01	10.44	5.68	5.83	6.43	**	+		
	HEMBB1000036	5.65	4.09	3.36	4.79	4.59	7.69	4.76	4.78	5.5				
	HEMBB1000037	6.62	4.31	5.17	7.83	6.16	9.26	6.18	5.41	5.32				
10	HEMBB1000039	3.3	1.35	2.08	5.56	6.46	6.46	3.88	3.39	2.84	**	+		
	HEMBB1000044	8.31	2.86	3	8.94	8.97	9.22	3.67	5.53	3.74				
	HEMBB1000048	4.16	1.72	3.61	5.69	6.15	8.14	3.51	4.43	3.25	*	+		
	HEMBB1000050	5.5	1.49	1.55	3.76	8.59	5.41	2.51	2.18	3.82				
	HEMBB1000054	5.55	2	2.53	9.07	6.03	8.7	7.15	3.88	5.66	*	+		
15	HEMBB1000055	24.4	16.2	17.8	18.24	19.34	22.83	9.69	8.54	9.54			*	-
	HEMBB1000059	8.8	6.35	7.84	16.75	19.27	21.09	9.69	10.78	9.65	**	+	*	+
	HEMBB1000072	9.51	4.64	5.32	12.83	10.68	11.19	7.97	7.6	5.64	*	+		
	HEMBB1000081	3.87	1.35	1.85	5.08	5.24	4.46	3.77	3.99	4.68	*	+		
	HEMBB1000083	4.74	2.08	3.56	8.88	6	6.36	3.2	5.07	6.07	*	+		
20	HEMBB1000089	3.6	2.1	3.13	10.31	7.12	8.77	3.62	4.07	4.02	**	+		
	HEMBB1000094	10.03	4.21	5.44	7.27	9.1	10.43	5.68	3.83	7.07				
	HEMBB1000097	2.21	1.8	1.66	3.6	3.78	2.43	2.31	1.65	1.94	*	+		
	HEMBB1000099	6	2.44	5.07	9.23	13.61	11.37	6.57	5.71	7.13	*	+		
	HEMBB1000103	11.08	5.29	6.37	9.34	10.14	10.72	4.69	6.24	4.67				
25	HEMBB1000106	6.42	4	5.39	8.37	6.27	6.82	6.5	5.47	4.71				
	HEMBB1000113	2.17	2	1.61	3.56	3.45	3.36	1.25	3.37	2.9	**	+		
	HEMBB1000119	4.55	2.45	4.15	5.3	3.89	4.98	2.17	5.09	5.65				
	HEMBB1000133	36.74	19.87	32.19	17.43	2.43	25.47	18.03	19.17	26.05				
	HEMBB1000134	8.1	5.02	4.94	5.99	6.85	11.63	3.4	5.64	6.33				
30	HEMBB1000136	4.52	2.17	1.45	2.82	2.31	2.54	3.01	2.62	4.93				
	HEMBB1000141	5.34	2.26	2.68	7.34	8.23	8.82	4.82	3.93	6.2	*	+		
	HEMBB1000144	4.28	3	3.58	12.18	6.95	9.35	4.11	4.95	6.86	*	+		
	HEMBB1000147	3	2.36	0.48	3.68	2.83	3.66	1.75	1.4	2.8				
	HEMBB1000152	4.26	2.59	2.98	3.85	2.52	3.5	2.62	3.23	3.16				
35	HEMBB1000154	3.63	1.65	1.97	5.05	4.98	5.15	2.28	3.46	4.23	*	+		
	HEMBB1000155	3.1	2.14	2.06	3.13	4.38	4.5	2.17	2.09	2.04	*	+		
	HEMBB1000173	11.42	5.05	6.29	19.61	16.74	17.56	10.24	8.45	9.62	**	+		
	HEMBB1000175	3.73	1.02	1.8	5.42	5.67	6.02	2.9	2.66	4.4	*	+		
	HEMBB1000176	5.82	2.57	3.52	6.79	7.3	6.93	5.44	4.12	6.38	*	+		
40	HEMBB1000198	2.93	1.33	0.9	2.24	0.81	1.87	1.77	0.77	1.87				
	HEMBB1000208	3.02	2.41	1.68	3.5	2.31	3.21	2.28	1.81	1.61				
	HEMBB1000209	4.47	2.11	2.26	5.05	5.4	5.79	2.1	3.16	2.24	*	+		
	HEMBB1000212	4.74	2.38	2.45	3.32	2.97	6.08	1.78	3.81	2.18				
	HEMBB1000215	12.22	6.74	7.81	16.21	19.51	21.21	10.04	11.3	10.31	*	+		
	HEMBB1000217	18.97	9.31	7.7	15.35	13.44	12.33	8.45	8.5	11.37				
45	HEMBB1000218	7.88	3.65	4.15	11.14	12.99	13.65	6.32	5.71	6.71	**	+		
	HEMBB1000226	9.75	5.82	3.67	9.36	7.18	7.09	5.55	5.63	5.67				
	HEMBB1000230	2.5	1.54	1.56	3.16	2.41	2.47	1.66	1.86	2.28				
	HEMBB1000240	2.54	1.04	1.59	2.21	2.34	2.83	1.25	2.23	1.86				
	HEMBB1000244	3.34	2.45	3.05	3.32	3.23	3.4	1.54	2.3	1.85		*	-	
50	HEMBB1000250	1.92	1.49	1.19	1.79	2.36	1.56	0.91	1.67	0.72				
	HEMBB1000258	8.84	4.21	4.45	9.29	10.64	11.84	4.49	4.85	5.22	*	+		
	HEMBB1000264	11.16	4.23	7.12	14.33	14.62	15.26	8.98	6.17	8.91	*	+		
	HEMBB1000266	7.49	4.1	3.58	5.54	5.59	6.27	4.11	3.08	5.07				
	HEMBB1000272	2.85	3.68	1.74	6.38	5.8	6.11	4.03	3.45	3.01	**	+		
55	HEMBB1000274	2.69	2.43	1.42	2.28	4.59	4.22	4.06	2.32	2.95				
	HEMBB1000276	2.16	0.94	0.86	1.1	3.12	1.78	0.56	0.79	1.49				
	HEMBB1000284	1.6	1.41	0.82	1.43	1.65	1.76	0.92	1.04	2.24				

Table 201

	HEMBA1007080	6.49	3.94	5.98	9.98	8.08	9.96	7.3	4.03	5.16	*	+		
	HEMBA1007084	6.15	4.73	3.3	7.53	11.43	12.96	4.4	6.54	5.84	*	+		
5	HEMBA1007085	11.57	6.03	6.42	14.47	16.1	16.28	10.38	7.67	10.37	*	+		
	HEMBA1007087	8.74	3.56	5.06	9.07	6.88	8.89	6.19	4.83	6.54				
	HEMBA1007089	4	1.08	1.76	3.78	3.23	3.34	2.04	2.11	2.69				
	HEMBA1007095	70.95	56.95	68.33	67.92	65.09	77.99	58.54	73.69	65.61				
	HEMBA1007101	8.13	4.48	3.34	8.09	6.98	8.85	8.22	8.26	10.62				
10	HEMBA1007104	5.96	2.89	2.91	5.64	4.63	4.55	3.31	2.95	4.74				
	HEMBA1007106	14.7	8.59	9.92	8.69	10.08	8.52	4.85	5.28	6.28		*	-	
	HEMBA1007112	2.54	1.7	2.5	2.24	3.01	4.69	1.68	1.95	2.11				
	HEMBA1007113	6.43	3.26	3.02	9.57	10.18	12	5.01	4.74	6.51	**	+		
	HEMBA1007121	15.29	6.28	8.37	20.01	26.55	24.61	14	11.02	13.68	*	+		
15	HEMBA1007129	4.97	2.15	2.01	4.97	3.24	4.17	3.35	2.12	2.39				
	HEMBA1007147	5.38	3.65	3.3	8.42	7.76	8.81	5.26	3.94	5.49	**	+		
	HEMBA1007149	4.94	2.77	3.26	6.72	8.54	6.33	4.3	4.3	4.5	*	+		
	HEMBA1007151	8.13	3.85	3.81	6.44	7.95	11.22	3.61	4.33	3.79				
	HEMBA1007172	7.56	3.48	4.13	7.44	5.05	7.28	4.12	5.13	4.04				
20	HEMBA1007174	5.89	2.49	3.67	3.93	3.83	5.88	2.72	4.89	2.91				
	HEMBA1007176	9.03	5.34	6.92	9.78	8.83	9.47	8.52	8	9.63				
	HEMBA1007178	32.55	18.88	15.14	19.06	21.65	17.59	9.08	8.47	7.89				
	HEMBA1007185	10.22	4.41	3.64	8.36	9.55	9.52	7.01	9.23	10.25				
	HEMBA1007186	5.79	5.42	2.99	5.38	6.38	4.34	5.55	4.63	4.28				
25	HEMBA1007194	10.77	5.25	6.27	6.05	8.58	8.52	4.54	4.6	4.86				
	HEMBA1007200	4.17	3	2.87	3.85	3.81	6.07	2.25	5.2	3.91				
	HEMBA1007203	7.33	3.38	4.4	6.6	5.9	7.76	3.26	5.38	5.31				
	HEMBA1007206	5.36	1.62	4.58	8.87	7.23	9.37	4.17	4.51	4.13	*	+		
	HEMBA1007224	4.31	3.41	3.02	7.21	8.94	7.9	5.84	2.98	5.52	**	+		
30	HEMBA1007226	8.11	2.53	3.92	5.1	4.65	5.56	3.57	3.89	3.99				
	HEMBA1007240	8.19	3.25	3.14	6.63	3.95	5.13	4.82	3.47	3.83				
	HEMBA1007241	2.29	1.82	2.1	4.38	3.16	4.31	2.93	3.05	2.77	*	+	**	+
	HEMBA1007242	3.53	1.89	1.63	1.79	3.06	2.59	1.23	2.17	2.42				
	HEMBA1007243	5.49	1.9	2.36	5.15	4.7	4.56	2.5	2.45	3.07				
35	HEMBA1007251	3.85	1.52	2.26	3.21	2.8	3.04	1.54	2.44	1.83				
	HEMBA1007256	2.11	1.7	2.58	4.85	3.63	4.4	1.15	2.23	2.07	**	+		
	HEMBA1007267	8.06	2.62	3.26	10.13	10.25	11.99	6.27	4.97	6.51	*	+		
	HEMBA1007273	2.76	1.75	1.08	1.92	1.89	2.71	1.52	1.78	1.02				
	HEMBA1007279	2.55	1.22	1.16	1.3	3.65	2.92	1.5	1.89	2.13				
40	HEMBA1007281	2.07	1.07	0.43	1.29	1.21	1.04	1.02	1.25	1.12				
	HEMBA1007283	6.62	2.63	3.23	3.75	3.81	4.38	2.75	2.29	3.75				
	HEMBA1007288	3.75	1.29	2.66	5.75	6.28	6.21	1.78	2.89	4.17	**	+		
	HEMBA1007291	3.22	0.96	1.72	2.4	3.14	3.81	1.55	2.4	4.26				
	HEMBA1007299	23.93	13.7	15.73	10.56	22.18	16.89	17.86	19.6	16.71				
45	HEMBA1007300	6.22	3.89	1.52	4.87	4.49	5.57	2.96	3.67	3.54				
	HEMBA1007301	4.77	2.47	2.12	3.91	6.06	4.53	5.31	3.92	4.11				
	HEMBA1007319	5.04	2.71	2.66	4.51	4.51	4.65	2.4	2.58	1.88				
	HEMBA1007320	3.5	1.62	1.5	3	2.95	3.58	2.72	2.88	2.98				
50	HEMBA1007322	28.33	24.69	28.25	30.89	47.79	40.83	20.16	16.66	16.95		**	-	
	HEMBA1007323	6.68	1.59	2.78	3.35	2.99	4.54	1.69	2.27	2.61				
	HEMBA1007326	16.87	9.35	13.09	29.82	36.45	31.07	12.34	13.22	15.57	**	+		
	HEMBA1007327	6.34	3.6	4.38	10.61	13.22	12.6	4.55	6.34	5.25	**	+		
	HEMBA1007332	13.26	4.92	5.19	6.74	8.15	8.34	6.2	5.28	6.24				
	HEMBA1007341	3.07	1.51	1.92	5.68	4.8	6.45	2.94	3.15	3.13	**	+		
	HEMBA1007342	3.54	1.8	1.84	3.52	2.33	2.69	2.06	2.55	1.53				
55	HEMBA1007347	6.86	4.49	4.81	9.76	12.67	13.86	6.9	5.92	8.38	**	+		
	HEMBA1007353	2.54	1.91	1.06	2.5	3.01	2.77	1.29	2.06	1.66				
	HEMBA1000085	5.95	3.76	2.97	7.43	7.91	9.69	2.81	4.53	3.98	*	+		

Table 200

	HEMBA1006849	7.06	2.5	3.59	4.52	8.98	7.67	3.87	4.24	3.66				
	HEMBA1006850	3.68	2.41	3.49	4.12	5.88	5.61	3.45	4.3	5.71	*	+		
5	HEMBA1006861	27.48	13.2	13.78	18.39	17.49	22.76	27.79	29.56	33.72				
	HEMBA1006865	7.81	4.59	4.59	10.66	9.55	9.31	6.64	6.59	6.33	*	+		
	HEMBA1006867	3.05	3.03	2.02	5.38	6.39	7.32	3.39	3.7	4.23	**	+		
	HEMBA1006873	3.17	1.82	1.33	4.27	2.94	4.49	4.02	3.19	3.84				
	HEMBA1006877	6.27	2.4	2.17	3.46	3.06	5.26	2.31	2.13	2.61				
10	HEMBA1006878	4.34	4.51	3.67	4.81	4.9	5.52	4.18	3.51	3.8				
	HEMBA1006879	17.53	11.84	12.53	14.59	8.5	17.01	9.97	13.17	14.21				
	HEMBA1006884	6.78	4.78	7.19	7.57	8.09	8	6.14	4.53	8				
	HEMBA1006885	14.47	10.91	10.29	11.14	13.59	13.12	8.92	9.99	11.51				
	HEMBA1006886	9.88	9.1	5.85	13.2	13.5	12.51	7.07	6.68	7.13	*	+		
15	HEMBA1006889	6.59	4.3	4.26	4.32	5.23	5.84	3.48	4.25	4.38				
	HEMBA1006896	16.57	11.14	9.96	13.58	12.6	17.46	13.28	8.89	13				
	HEMBA1006900	11.28	4.72	4.94	6.26	7.37	10.33	5.94	4.33	6.61				
	HEMBA1006902	2.57	1.63	2.97	3.06	2.5	3.31	2.84	4.11	2.62				
	HEMBA1006912	9.86	3.49	5.48	8.69	10.41	10.91	5.79	6.47	5.76				
20	HEMBA1006914	14.14	7.94	10.37	14.19	14.05	16.96	6.19	5.9	9.72				
	HEMBA1006916	9.91	7.1	4.15	7.61	7.72	7.02	3.84	3.67	4.33				
	HEMBA1006921	5.33	2.22	1.77	2.75	3.09	2.98	2.63	2.52	3.45				
	HEMBA1006926	4.69	3.93	4.04	8.12	6.37	6.61	5.19	4.08	4.59	**	+		
	HEMBA1006927	2.56	1.45	1.11	4.26	3.27	5.93	2.47	2.76	2.3	*	+		
25	HEMBA1006929	3.54	1.38	2	3	2.51	2.71	2.05	2.99	1.84				
	HEMBA1006936	6.81	2.92	3.95	7.43	7.48	8.89	3.83	5.74	3.63				
	HEMBA1006938	1.33	0.26	0.47	5.31	1.59	1.56	1.54	1.69	0.91				
	HEMBA1006941	16.53	11.05	11.6	12.22	7.8	9.63	10.28	8.93	11.52				
	HEMBA1006942	8.19	4.07	6.53	8.73	9.65	14.5	10.35	7.57	10.44				
30	HEMBA1006945	25.04	16.05	14.06	21.51	28.59	29.47	11.94	11.2	11.54				
	HEMBA1006949	2.9	1.1	0.96	1.63	1.82	4.13	0.8	1.36	1.9				
	HEMBA1006952	3.78	1.55	1.57	2.91	2.65	3.54	2.84	4.46	4.01				
	HEMBA1006960	10.85	6.07	5.14	11.23	9.86	8.27	10.08	9.22	8.03				
	HEMBA1006973	3.3	3.69	3.3	7.1	4.93	5.77	3.56	4.84	3.61	*	+		
	HEMBA1006974	5.62	2.6	4.96	7.66	9.22	8.05	3.96	5.98	3.51	*	+		
35	HEMBA1006976	2.71	1.15	1.73	3.59	2.62	4.04	2.12	3.56	2.05				
	HEMBA1006989	0.83	0.32	0.23	0.34	1.18	1.21	0.38	0.32	1.18				
	HEMBA1006993	7.77	3.49	2.52	13.12	7.8	8.64	3.93	4.49	6.13				
	HEMBA1006996	1.18	0.27	0.63	0.83	0.78	0.99	0.66	1	0.65				
40	HEMBA1007001	5.49	3.33	4.13	8.5	12.04	10.88	5.67	5.2	5.4	**	+		
	HEMBA1007002	5.81	2.2	3.66	4.91	3.97	4.41	3.34	3.47	2.91				
	HEMBA1007013	3.72	1.85	2.69	3.52	4.62	4.75	3.38	4.47	2.43				
	HEMBA1007016	3.01	1.36	1.4	2.83	2.49	3.86	1.86	2.87	2.52				
	HEMBA1007017	0.36	0.56	0.46	1.7	1.63	2.47	0.33	1.63	0.45	**	+		
	HEMBA1007018	9.21	6.01	5.67	4.76	4.66	5.23	4.98	4.73	5.19				
45	HEMBA1007044	9.95	5.07	6.68	9.58	7.21	9.83	7.74	6.77	8.84				
	HEMBA1007045	2.71	0.74	1.32	2.37	2.16	3.02	2.18	2.14	3.63				
	HEMBA1007051	4.5	1.5	2.49	3.07	5.64	4.56	4.1	1.96	3.57				
	HEMBA1007052	2.79	1.47	1.81	2.94	3.23	3.76	2.14	1.66	1.93	*	+		
	HEMBA1007053	2.08	1.3	1.15	3.49	2.13	3.23	2.17	2.69	2.75	*	+	*	+
50	HEMBA1007057	4.25	1.9	2.27	4.24	3.69	4.46	2.89	2.65	3.08				
	HEMBA1007062	6.55	4.08	2.49	3.19	4.21	5.74	3.61	4.05	2.65				
	HEMBA1007063	7.3	3.36	3.27	9.41	8.17	9.36	5.6	4.6	6.77	*	+		
	HEMBA1007066	4.89	2.13	1.75	9.06	3.28	4.77	3.8	2.51	6.25				
	HEMBA1007069	3.01	1.67	1.08	4.66	3.81	6.58	2.68	2.84	4.17	*	+		
55	HEMBA1007073	3.81	1.52	1.06	5.19	3.51	7.75	1.69	2.27	1.95				
	HEMBA1007076	8.06	4.01	4.39	8.06	8.27	7.87	6.08	3.95	7.1				
	HEMBA1007078	44.29	25.47	26.67	39.89	48.08	43.86	17.13	16.51	23.74				

Table 199

	HEMBA1006595	4.6	1.32	2.47	6.45	3.43	5.48	2.48	3.17	3.35			
5	HEMBA1006597	6.19	2.47	4	9.61	11.89	11.02	4.43	7.63	6.03	**	+	
	HEMBA1006606	5.22	2.34	3.15	5.5	7.09	8.72	4.17	3.59	5.67	*	+	
	HEMBA1006612	5.88	3.13	2.66	9.51	7.07	8.75	4.24	4.07	8.12	*	+	
	HEMBA1006617	6.23	2.4	3.25	7.51	8.15	9.4	4.22	3.47	5.72	*	+	
	HEMBA1006624	21.51	11.59	11.39	8.91	10.89	11.11	15.72	17.22	19.01			
	HEMBA1006631	11.14	7.16	5.63	14.71	13.36	15.13	9.17	8.76	9.27	*	+	
10	HEMBA1006635	3.5	1.48	1.8	6.1	5.02	7.12	3.2	2.77	3.44	*	+	
	HEMBA1006639	5.83	1.94	3.55	4.08	4.21	4.37	3.14	4	3.07			
	HEMBA1006643	8.1	3.39	6.04	7.92	5.21	8.41	3.69	6.07	4.57			
	HEMBA1006648	7.17	4.23	2.23	4.85	5.86	6.95	5.26	5.13	6.25			
	HEMBA1006652	7.55	5.4	7.95	14.31	13.73	13.23	6.43	7.1	11.54	**	+	
15	HEMBA1006653	6.97	4.5	3.06	4.22	5.74	4.88	4.94	3.37	4.63			
	HEMBA1006658	7.71	4.81	3.99	9.26	8.5	11.38	5.3	4.42	6.47	*	+	
	HEMBA1006659	7.41	4.7	3.7	5.26	4.56	4.46	6.04	3.81	4.25			
	HEMBA1006665	1.62	1.53	0.92	2.6	1.66	1.94	1.6	1.36	2.14			
	HEMBA1006666	2.8	1.45	1.19	5.48	2.51	3.57	1.85	1.35	3.75			
20	HEMBA1006671	4.48	2.13	2.48	3.04	6.4	6.86	3.55	4.19	4.16			
	HEMBA1006674	4.97	3.16	4.4	5.76	5.14	7.87	4.61	3.42	4.54			
	HEMBA1006676	10.46	5.08	3.85	9.54	8.88	9.7	6.21	4.55	6.44			
	HEMBA1006682	2.27	1.69	1.34	3.17	2.06	2.05	4.61	1.08	3.99			
	HEMBA1006688	6.01	4.37	2.5	5.47	6.02	6.19	4.31	2.6	4.14			
25	HEMBA1006695	4.5	1.72	1.74	6.75	6.52	5.65	3.76	2.82	3.65	*	+	
	HEMBA1006696	12.87	6.14	7.8	9.63	11.85	11.77	5.03	6.37	5.41			
	HEMBA1006702	2.64	1.17	1.68	3.05	1.99	2.26	2.52	2.64	2.72			
	HEMBA1006707	6.85	2.92	3.19	5.67	3.46	4.24	2.84	4.21	4.09			
	HEMBA1006708	8.39	4.87	3.01	5.26	5	6.1	6.53	3.85	5.31			
30	HEMBA1006709	6.65	3.16	3.47	4.07	5.63	4.68	6.45	3.52	4.44			
	HEMBA1006717	8.88	2.4	4.14	4.44	3.37	2.93	4.5	3.69	4.56			
	HEMBA1006724	3.81	3.86	1.52	3.61	3.98	3.44	2.83	2.11	3.27			
	HEMBA1006731	7.51	3.16	2.94	4.8	6.48	6.17	3.61	3.73	4.13			
	HEMBA1006737	5.15	2.61	1.58	2.17	3.41	5.22	2.11	2.54	2.79			
35	HEMBA1006742	4.81	2.29	1.84	6.06	4.83	6.03	2.78	3.29	3.24			
	HEMBA1006743	7.87	4.47	4.75	8.29	5.08	7.45	3.49	6.04	3.57			
	HEMBA1006744	10.08	3.77	3.8	14.22	11.75	16.16	7.99	6.73	6.12	*	+	
	HEMBA1006749	3.53	3.65	2.98	4.2	4.74	5.34	4.08	3.16	4.37	*	+	
	HEMBA1006752	23.27	11.82	13.93	14.5	12.58	14.16	12.27	10.32	10.17			
40	HEMBA1006754	1.86	1.19	1.02	4.17	4.31	3.82	2.65	2.7	3.64	**	+	+
	HEMBA1006758	8.94	5.63	3.63	4.07	5.41	4.85	3.57	4.16	2.9			
	HEMBA1006767	3.06	1	1.61	2.61	3	3.72	1.69	2.22	2.41			
	HEMBA1006770	13.78	5	6.03	7.89	10.06	11.16	4.74	6.16	6.66			
	HEMBA1006779	10.4	3.74	5.54	13.72	14.85	14.12	6.44	8.3	7.48	*	+	
	HEMBA1006780	7.08	3.47	3.59	13.82	10.5	10.84	7.75	5.05	6.84	*	+	
45	HEMBA1006789	4.72	5.04	4.21	3.76	5.14	4.99	3.13	3.89	4.1			
	HEMBA1006795	8.9	4.61	4.5	12.12	13.21	10.55	5.76	4.72	5.99	*	+	
	HEMBA1006796	7.65	2.94	3.34	4.85	4.95	4.32	4.94	2.99	4.97			
	HEMBA1006805	6.94	4.11	2.79	6.38	6.88	9.56	5.8	4.23	4.32			
	HEMBA1006807	41.87	16.77	24.31	34.14	30.28	35.28	17.64	15.46	18.76			
50	HEMBA1006813	2.76	1.69	0.75	4.56	3.37	2.07	1.6	1.54	1.68			
	HEMBA1006819	5.85	2.89	4.93	3.44	4.06	3.24	2.46	2.97	2.69			
	HEMBA1006821	4.19	2.43	1.27	6.45	6.7	7.35	2.91	3.28	3.89	**	+	
	HEMBA1006824	6.62	2.68	2.84	7.05	7.56	7.49	5.6	3.8	3.89			
	HEMBA1006832	34.7	31.52	23.59	34.4	30.12	38.56	20.42	16.73	24.06			
	HEMBA1006834	23.99	11.25	13	13.06	14.6	15.62	11.33	8.76	11.95			
55	HEMBA1006835	4	1.43	1.9	3.69	3.2	3.36	2.22	2.96	1.96			
	HEMBA1006843	103.5	33.5	66.05	133.5	136.8	126.1	52.72	23.1	39.64	*	+	

Table 198

	HEMBA1006360	7.98	4.95	5.62	5.47	3.56	4.4	1.94	2.36	2.5		*	-
	HEMBA1006364	3.11	1.13	2.29	5.13	3.18	5.17	4.53	10.75	7.49		*	+
5	HEMBA1006377	9.83	4.08	4.81	9.68	6.73	11.12	5.12	6.15	6.04			
	HEMBA1006380	8.33	2.76	3.16	7.63	7.47	9.64	4.68	4.63	5.81			
	HEMBA1006381	27.84	15.11	15.63	23.73	22.47	28.24	18.48	12.8	18.34			
	HEMBA1006385	9	3.81	3.21	10.12	10.82	10.06	5.78	6.06	7.86			
	HEMBA1006390	10.59	5.3	6.11	5.74	8.16	9.4	6.45	5.63	5.84			
10	HEMBA1006391	5.9	2.52	2.93	4.19	2.98	3.66	3.7	3.39	4.92			
	HEMBA1006398	1.24	0.85	0.78	1.46	2.48	2.33	1.32	1.57	1.25	*	+	
	HEMBA1006405	6.46	2.31	3.39	3.97	5.97	7.86	4.98	4.43	6.01			
	HEMBA1006410	10.66	4.34	6.26	48.24	9.18	6.95	5.67	6.99	5.47			
	HEMBA1006416	7.58	3.75	4.83	11.17	11.6	10.4	5.86	5.53	5.52	**	+	
15	HEMBA1006418	4.85	2.81	2.36	4.42	4.54	5.46	2.95	3.19	4.26			
	HEMBA1006419	8.31	4.08	4.44	13	13.16	12.95	7.56	6.59	6.8	**	+	
	HEMBA1006421	2.57	1.36	2.21	4.58	3.93	3.93	2.69	2.86	2.95	**	+	
	HEMBA1006424	1.92	1.1	0.54	1.6	1.43	2	1.13	1.24	1.46			
	HEMBA1006426	6.91	3.24	3.97	14.78	13.77	12.89	6.5	5.72	7.38	**	+	
20	HEMBA1006430	4.14	1.54	1.15	3.22	4.8	4.46	2.25	2.55	3.21			
	HEMBA1006438	3.24	1.25	2.86	4.15	5.58	5.24	2.63	2.65	2.43	*	+	
	HEMBA1006445	5.47	3.56	1.09	4.34	6.2	5.79	5.24	5.14	9.95			
	HEMBA1006446	2.47	0.4	0.6	1.78	0.97	2.17	2.61	1.77	0.98			
	HEMBA1006456	9.3	7.18	5.88	27.97	39.53	36.06	25.26	23.55	25.96	**	+	**
25	HEMBA1006461	3.9	2.47	2.09	3.96	6.32	5.5	3.18	2.5	2.97			
	HEMBA1006467	3.36	2.3	2.41	1.89	3.11	2.94	1.06	2.01	1.22		*	-
	HEMBA1006470	3.32	2.6	1.74	4.73	4.89	6.17	2.71	2.99	2.35	*	+	
	HEMBA1006471	2.77	2.01	2.5	2.54	4.17	4.09	1.83	2.8	1.93			
	HEMBA1006474	3.4	0.88	1.69	1.95	2.26	1.5	0.73	1.98	1.64			
30	HEMBA1006476	7.63	2.81	3.49	7.03	6.55	10.28	5.71	6.01	8.9			
	HEMBA1006482	53.61	36.99	43.8	47.46	64.27	63.44	24.67	21.43	26.34		*	-
	HEMBA1006483	5.77	3.34	3.12	9.27	6.33	10.42	4.67	4.49	5.8	*	+	
	HEMBA1006485	2.4	0.96	1.41	4.2	4.91	5.55	9.43	7.34	8.87	**	+	**
	HEMBA1006486	22.07	14.47	14.17	13.5	21.65	20.32	9.55	5.18	8.79		*	-
35	HEMBA1006489	2.84	0.31	0.23	0.65	1.22	0.91	1.3	1.95	0.72			
	HEMBA1006492	22.55	16.4	18.02	18.63	19.03	19.21	4.75	5.92	5.79		**	-
	HEMBA1006494	1.6	0.13	1.42	1.49	1.22	1.56	0.94	0.97	0.8			
	HEMBA1006497	4.42	2.46	1.3	2.7	3.38	4.13	3.19	2.22	2.93			
	HEMBA1006501	6.77	2.17	3.41	4.37	3.72	6.05	2.94	2.94	4.13			
40	HEMBA1006502	14.3	11.26	8.46	15.96	17.52	16.95	15.96	11.43	17.31	*	+	
	HEMBA1006507	3.4	0.73	1.23	5.85	4.08	5.84	2.92	3.88	4.16	*	+	
	HEMBA1006517	4.63	2.62	2.31	5.72	6.14	5.82	3.68	4.27	4.71	*	+	
	HEMBA1006521	3.02	1.72	1.98	2.24	2.27	2.97	3.41	3.45	2.36			
	HEMBA1006529	6.54	5.38	7.96	6.72	7.42	7.81	5.9	6.56	6.87			
	HEMBA1006530	1.54	0.77	2.01	2.93	1.8	2.4	1.35	1.69	1.44			
45	HEMBA1006535	2.61	2.15	0.64	3.13	3.63	3.67	2.05	1.48	2.17			
	HEMBA1006536	5.93	3.85	4.16	6.52	8.47	8.22	4.62	4.48	4.48	*	+	
	HEMBA1006540	4.27	2.17	1.9	4.22	2.42	3.65	2.42	2.05	2.05			
	HEMBA1006544	1.52	0.67	1.46	2.15	3.36	3.6	2.21	2.99	2.6	*	+	*
	HEMBA1006546	4.48	4.88	3.24	16.24	9.73	11.7	5.09	6.41	8.5	*	+	
50	HEMBA1006549	2.11	0.58	0.86	2.8	1.88	2.9	1.86	1.87	1.42			
	HEMBA1006559	5.16	2.1	4.6	12.73	9.91	12.55	8.17	8.31	7.9	**	+	*
	HEMBA1006562	2.22	0.76	1.85	3.22	2.69	2.84	1.63	3.25	2.16	*	+	
	HEMBA1006566	1.5	1.62	0.13	0.8	1.28	0.97	1.14	1.33	0.88			
	HEMBA1006569	4.26	2.46	1.96	4.02	5.76	5.28	3.58	2.64	3.6			
55	HEMBA1006572	1.59	0.24	0.54	0.56	0.89	1.09	1.05	1.04	1.21			
	HEMBA1006579	2.51	1.31	1.43	2.63	2.93	3.26	6.37	6.01	6.92	*	+	**
	HEMBA1006583	3.62	1.64	2.4	3.61	3.89	4.77	3.38	3.77	2.05			+

Table 197

	HEMBA1006108	5.03	2.45	2.82	5.62	4.96	3.72	3.28	3.95	3.28			
	HEMBA1006114	5.25	4.63	5.08	7.3	10.42	7.17	4.76	5.44	5.87	*	+	
5	HEMBA1006121	6.32	2.33	4.31	5.84	6.44	7.33	4.17	6.55	4.7			
	HEMBA1006124	3.12	2.28	2.5	3.33	4.9	2.3	1.89	3.9	2.53			
	HEMBA1006125	10.14	8.44	4.52	7.52	17.2	16.18	9.52	10.87	14.31			
	HEMBA1006130	2.62	2.68	2.39	2.72	3.08	4.43	3.7	4.3	4.1		**	+
	HEMBA1006138	7.26	4.73	3.72	9.3	11.39	10.14	5.49	5.98	7.37	*	+	
10	HEMBA1006142	6.22	3.63	4.24	7.33	10.18	10.72	6.57	6.34	6.19	*	+	
	HEMBA1006150	16.28	10.88	9.66	15.57	15.3	13.33	6.57	7.84	7.68			
	HEMBA1006151	8.94	6.23	8.3	9.44	9.41	9.8	14.8	13.36	17.11		**	+
	HEMBA1006155	4.31	2.12	3.11	2.99	2.19	2.62	2.75	4.44	3.92			
	HEMBA1006158	1.99	2.23	1	5.52	2.28	1.62	0.79	3.02	2.04			
15	HEMBA1006164	7.82	6.93	4.48	10.95	14.83	12	6.46	6.96	7.98	*	+	
	HEMBA1006171	3.78	1.96	1.78	2.93	3.7	4.2	6.07	5.07	5.46		*	+
	HEMBA1006173	3.13	1.34	2.45	2.99	4.82	4.35	2.87	4.45	2.71			
	HEMBA1006176	17.29	15.19	12.08	17.72	24.16	22.1	76.2	63.22	78.98		**	+
	HEMBA1006182	2.42	1.06	1.52	2.8	3.22	2.43	1.16	3.5	1.94			
20	HEMBA1006197	6.41	5.46	4.82	12.32	9.66	9.7	4.32	5.89	5.15	**	+	
	HEMBA1006198	9.58	7.2	6.52	9.4	9.55	10.32	5.65	8.56	6.79			
	HEMBA1006213	2.56	0.9	1.99	3.02	4.19	4.18	1.76	2.58	3.01	*	+	
	HEMBA1006217	23.81	12.95	14.09	28.71	29.21	22.65	54.8	57.77	74.75		**	+
	HEMBA1006226	45.81	48.81	55.06	71.05	67.87	69.04	34.7	30.76	48.77	**	+	
25	HEMBA1006235	2.69	1.66	2.93	2.89	2.63	3.42	3.26	2	2.73			
	HEMBA1006248	4.57	1.66	2.14	4.47	3.25	4.51	3.57	3.35	2.98			
	HEMBA1006251	7.31	5.13	5.62	8.77	8.46	10.53	8.03	7.68	7.92	*	+	*
	HEMBA1006252	2.83	2.65	0.76	1.86	2.33	3.7	2.51	1.94	2.08			
	HEMBA1006253	5.52	3.08	3.71	4.06	4.47	4.75	2.99	2.68	1.89			
	HEMBA1006259	4.17	1.88	2.86	4.37	4.88	6.45	2.66	2.31	3.49			
30	HEMBA1006261	6.4	3.95	3.39	6.02	5.83	6.2	5.45	3.63	10.61			
	HEMBA1006268	3.66	2.08	1.88	4.46	4.9	5.18	2.58	2.36	4.27	*	+	
	HEMBA1006271	7.71	2.93	4.51	11.62	12.09	12.3	7.07	5.33	10.91	**	+	
	HEMBA1006272	2.81	1.63	1	2.86	2.92	3.49	2.16	1.96	2.4			
	HEMBA1006273	5.39	2.09	3.07	4.81	3.79	4.4	5.32	3.06	3.91			
35	HEMBA1006276	2.93	1.9	3.24	3.4	4.55	3.76	2.55	1.66	2.29			
	HEMBA1006278	1.93	1.63	1.33	4.06	4.19	3.8	2.43	1.58	2.09	**	+	
	HEMBA1006283	7.35	3.25	3.5	4.82	5.8	5.93	4.92	3.12	4.11			
	HEMBA1006284	3.83	2.26	2.04	5.58	2.8	4.34	3.15	2.33	3.82			
	HEMBA1006291	4.96	1.36	1.34	4.1	2.68	4.41	3.86	3.13	3.18			
40	HEMBA1006292	2.77	2.02	1.73	2.32	2.22	1.89	2.26	1.67	2.38			
	HEMBA1006293	3.02	0.92	0.7	1.9	1.76	2.36	1.54	1.85	1.56			
	HEMBA1006299	3.49	2.22	1.51	13.99	12.93	16.92	7.99	7.28	10.15	**	+	**
	HEMBA1006309	5.39	3.08	3.38	5.38	6.85	7.74	3.06	4.11	4.45			
	HEMBA1006310	3.7	2.35	2.24	5.29	3.06	3.56	2.59	4.56	4.32			
45	HEMBA1006311	8.15	4.04	4.72	3.8	4.97	7.43	4.03	5.26	5.64			
	HEMBA1006313	2.58	0.57	1	1.55	1.73	1.85	2.63	1.09	1.3			
	HEMBA1006316	2.99	1.66	1.44	1.74	2.62	2.14	2.59	1.79	1.84			
	HEMBA1006328	4.68	2.1	1.68	6.39	5.95	6.83	4.27	3.72	3.95	*	+	
	HEMBA1006334	2.26	1.44	1.07	1.93	1.34	1.2	1.12	1.33	0.99			
50	HEMBA1006335	10.13	6.95	5.67	4.72	4.51	6.4	10.88	11.65	14.01		*	+
	HEMBA1006344	4.43	2.82	4.27	9.97	8.14	7.72	4.65	6.26	4.98	**	+	
	HEMBA1006347	5.25	2.13	2.64	4.75	3.92	6.02	3.02	3.83	3.69			
	HEMBA1006349	6.07	2.73	2.89	4.44	4.96	6.67	4.94	4.8	4.22			
	HEMBA1006352	3.21	2.07	2.2	4.23	3.53	3.79	3.57	2.77	2.87	*	+	
55	HEMBA1006357	9.36	4.79	5.03	14.77	13.42	14.23	7.21	5.46	6.81	**	+	
	HEMBA1006358	4.06	2.27	1.93	3.39	4.53	4.11	2.56	2.11	2.8			
	HEMBA1006359	11.9	9.22	8.59	18.27	21.46	21.84	9.68	5.92	7.59	**	+	

Table 196

	HEMBA1005829	7.71	4.11	4.16	9.68	9.82	10.65	5.68	6.05	6.18	*	+		
5	HEMBA1005833	5.58	4.05	3.69	5.07	5.16	5.6	4.09	4.46	5.21				
	HEMBA1005834	6.55	4.34	5.21	12.06	12.18	15.25	4.16	7.19	5.66	**	+		
	HEMBA1005844	55.19	32.63	42.62	52.31	50.88	44.4	13.71	22.39	16.54		*	-	
	HEMBA1005852	14.32	7.35	8.88	11.42	13.87	12.28	12.12	9.6	10.71				
	HEMBA1005853	4.46	3.87	2.7	5.48	7.15	7.24	6.76	3.1	4.03	*	+		
10	HEMBA1005878	10.9	9.31	6.82	15.29	18.75	18.35	11.26	9.02	9.91	**	+		
	HEMBA1005883	2.8	3.02	2.09	2.99	4.75	3.12	3.03	3.43	2.83				
	HEMBA1005884	1.78	1.18	0.5	2.41	2.22	1.91	2.16	1.93	1.73				
	HEMBA1005891	1.55	1.14	0.52	2.25	4.37	4.09	2.08	2.69	1.69	*	+		
	HEMBA1005894	3.43	2.12	2.97	5.44	5.86	5.44	2.54	4.52	2.77	**	+		
15	HEMBA1005898	16.67	8.8	11.51	11.61	18.53	21.97	6.97	12.21	8.22				
	HEMBA1005902	4.41	3.46	2.55	2.97	3.31	3.57	3.63	4.8	4.43				
	HEMBA1005907	1.14	1	0.32	1.39	1.9	1.41	1.83	2.17	1.38		*	+	
	HEMBA1005909	0.96	0.99	0.06	0.74	1.52	0.83	1.8	0.82	0.95				
	HEMBA1005911	5.56	3.24	3.54	5.59	8.12	8.18	4.97	3.97	5.62	*	+		
	HEMBA1005912	6.61	6.28	5.64	8.63	10.33	8.51	7.27	7.15	4.9	*	+		
20	HEMBA1005913	3.32	1.87	2.67	4.85	5.83	5.39	4.23	6.09	5.19	**	+	*	+
	HEMBA1005921	5.08	3.6	4.07	7.96	11.09	11.08	3.93	6.12	4.64	**	+		
	HEMBA1005922	9.29	4.86	8.75	10.31	11.79	14.59	5.42	7.95	6.59				
	HEMBA1005929	9.26	6.15	5.27	8.35	12.25	12.51	8.91	7.98	6.88				
25	HEMBA1005931	13.37	8.03	6.05	13.2	15.89	16.14	10.01	9.04	10.17				
	HEMBA1005934	11.83	7.65	6.91	11.33	21.92	13.8	6.94	9.42	10.1				
	HEMBA1005945	9.41	6.42	4.64	6.1	7.01	8.67	8.01	6.77	7.06				
	HEMBA1005962	2.52	1.69	1.85	2.52	2.44	3.11	1.69	3.18	2.61				
	HEMBA1005963	1.58	1.29	0.83	2.22	2.32	1.65	0.75	2.23	1.58				
	HEMBA1005990	53.63	37.05	35.87	22.88	28.11	30.49	25.75	38.21	38.5				
30	HEMBA1005991	4.36	2.88	2.52	7.83	8.53	8.07	3.66	3.18	4.37	**	+		
	HEMBA1005999	7.25	4.04	3.51	7.81	9.22	8.54	5.71	6.17	5.07	*	+		
	HEMBA1006002	4.03	2.6	1.83	2.32	2.41	2.99	3.56	4.2	3.68				
	HEMBA1006005	3.58	3.7	2.47	1.41	2.98	2.78	2.19	3.32	3.16				
	HEMBA1006011	28.82	13.22	19.62	6.69	8.42	8.26	9.43	7.34	8.25	*	-		
35	HEMBA1006013	4.9	3.69	2.44	2.82	3.64	2.69	3.14	3.46	2.63				
	HEMBA1006016	5.42	2.01	3.02	4.73	5.78	5.82	3.09	4.11	3.71				
	HEMBA1006019	4.75	3.24	2.19	2.66	6.4	5.83	2.01	3.58	3.27				
	HEMBA1006021	5.17	2.64	3.76	13.9	20.33	23.22	9.49	12.71	9.39	**	+	**	+
	HEMBA1006022	6.7	7.43	3.24	7.5	7.39	6.93	5.83	6.01	8.3				
40	HEMBA1006031	4.39	5.2	2.1	3.55	7.12	4.25	2.82	4.39	3.34				
	HEMBA1006035	3.57	1.83	2.1	2.68	3.31	3.32	3.52	3.36	3.1				
	HEMBA1006036	11.47	5.72	5.91	13.84	22.61	19.36	7.96	7.38	10.66	*	+		
	HEMBA1006042	5.24	3.69	2.84	6.48	8.01	7.56	4.36	7.77	4.18	*	+		
	HEMBA1006044	1.69	0.79	0.7	2	1.1	1.58	0.9	2.05	1.25				
45	HEMBA1006045	4.3	3.06	2.36	5.33	6.87	5.75	4.69	7.34	3.91	*	+		
	HEMBA1006048	5.42	3.01	4.33	5.37	6.23	4.19	3.1	3.81	2.5				
	HEMBA1006053	5.79	4.06	2.48	4.5	6.49	3.55	3.66	3.74	4.34				
	HEMBA1006055	1.82	1.84	1.28	1.8	2.36	2.19	1.75	2.52	2.48				
	HEMBA1006058	4.72	2.18	2.21	2.56	3.95	3.04	3.54	3.28	3.39				
	HEMBA1006063	15.52	11.99	10.03	16.08	16.03	13.94	13.46	9.12	10.83				
50	HEMBA1006067	1.98	1.55	1.25	1.72	2.65	1.7	2.65	2.72	2.71		**	+	
	HEMBA1006081	3.98	3.25	2.94	3.52	4.19	3.86	2.74	3.6	2.18				
	HEMBA1006089	10.88	7.08	9.01	8.5	7.48	9.62	5.26	4.65	6.13	*	-		
	HEMBA1006090	2.72	1.74	2.31	2.48	4.09	2.53	1.71	3.25	2.66				
	HEMBA1006091	8.41	4.97	5.38	6.67	13.08	9.53	6.54	5.82	7.45				
55	HEMBA1006093	4.66	3.46	1.8	4.22	4.68	5.91	4.02	4.39	5.59				
	HEMBA1006099	8.2	2.83	3.57	7.25	7.27	6.62	8.7	6.8	7.75				
	HEMBA1006100	4.94	3.58	3.48	6.75	7.94	7.84	5.43	4.52	4.63	**	+		

Table 195

	HEMBA1005548	2.54	2.07	2.02	3.97	6.52	4.14	3.37	3.9	3.32	*	+	**	+
5	HEMBA1005552	9.98	4.38	5.49	14.16	16.16	16.24	6.88	9.1	7.91	**	+		
	HEMBA1005558	5.62	4.78	4.01	4.12	4.94	4.94	2.89	4.54	2.98				
	HEMBA1005568	4.56	2.35	2.64	4.41	6.84	7.67	2.66	3.77	3.75				
	HEMBA1005570	22.81	14.72	12.89	3.4	5.87	4.67	2.86	3.28	4.18	*	-	*	-
	HEMBA1005576	3.57	2.9	1.76	5.63	4.9	6.27	3.31	4.43	3.65	*	+		
10	HEMBA1005577	3.28	1.8	1.85	2.52	3.76	3.29	1.78	2.45	2.1				
	HEMBA1005581	6.44	3.47	3.35	11.86	10.8	9.38	9.31	8.35	7.77	**	+	*	+
	HEMBA1005582	3.79	2.19	1.67	4.94	4.83	5.37	3.11	3.69	2.69	*	+		
	HEMBA1005583	2.18	2.16	1.54	2.99	3.77	4.66	2.3	2.75	1.62	*	+		
	HEMBA1005588	3.6	2.49	3.31	8.28	7.89	9.86	3.63	5.17	4.67	**	+		
	HEMBA1005593	3.44	3.2	2.65	4.18	6.03	3.87	2.97	3.28	2.95				
15	HEMBA1005595	2.58	2.31	1.83	3.46	4.98	5.89	3.2	2.15	3.87	*	+		
	HEMBA1005597	13.38	9.58	8.44	10.53	12.2	11.02	8.53	9.47	8.93				
	HEMBA1005606	12.27	7.53	6.44	5.89	7.11	6.3	8.22	8.78	11.95				
	HEMBA1005609	5.25	3.66	3.27	10.52	11.83	10.56	4.85	5.36	5.71	**	+		
	HEMBA1005616	5.15	3.24	2.69	6.77	7.27	7.69	4.76	5.32	4.74	*	+		
20	HEMBA1005621	5.71	4.59	4.34	4.48	5.05	3.45	2.83	4.75	2.83				
	HEMBA1005627	4.83	2.61	2.82	6.51	8.02	6.48	3.29	4.97	4.83	*	+		
	HEMBA1005628	5.64	3.83	3.44	12.81	11.82	14.97	10.64	9.94	13.34	**	+	**	+
	HEMBA1005631	2.21	1.39	0.65	2.83	4.04	3.15	5.61	3.11	3.88	*	+	*	+
	HEMBA1005632	11.01	3.49	3.42	8.83	9.02	7.82	5.06	4.35	5.44				
25	HEMBA1005634	6.35	2.76	2.05	5.36	8.63	6.5	4.98	5	6.87				
	HEMBA1005662	1.07	1.53	1.02	2.26	2.43	2.33	2.04	1.73	1.38	**	+		
	HEMBA1005666	4.52	3.82	4.32	9.91	8.09	7.3	6.48	6.28	5.06	**	+	*	+
	HEMBA1005670	2.29	2.27	1.9	7.3	6.51	7	3.1	7.04	3.71	**	+		
	HEMBA1005671	3.97	1.07	3.6	3.68	3.22	2.26	4.53	6.9	3.6				
30	HEMBA1005679	4.26	2.11	3.13	6.55	7.51	6.35	2.51	4.92	3.8	**	+		
	HEMBA1005680	6.79	3.09	2.88	6.98	9.15	8.11	7.19	3.45	6.54				
	HEMBA1005685	5.15	2.24	2.86	3.16	3.75	6.06	3.75	2.67	3.13				
	HEMBA1005698	6.46	4.64	3.65	6.51	6.49	8.04	4.48	5.97	6.27				
	HEMBA1005699	2.04	1.37	1.03	2.33	2.8	2.44	1.39	3.16	0.93	*	+		
35	HEMBA1005703	1.57	1.14	0.53	2.63	1.8	1.22	0.95	3.02	1.71				
	HEMBA1005705	4.78	2.62	3.65	8.55	5.59	7.85	3.94	5.46	2.65	*	+		
	HEMBA1005712	1.7	0.73	0.42	2.78	2.29	2.36	1.03	2.79	1.13	*	+		
	HEMBA1005717	1.99	1.9	1.57	4.59	18.53	4.07	1.65	3.65	2.24				
	HEMBA1005718	12.46	6.17	5.4	10.4	11.53	8.97	6.74	7.19	8.25				
40	HEMBA1005721	15.4	8.95	6.41	11.18	12.64	11.59	11.3	10.89	13.73				
	HEMBA1005722	11.88	7.25	5.73	15.89	16.63	13.24	10.07	13.96	12.55	*	+		
	HEMBA1005724	4.23	1.39	1.12	1.47	3.11	2.3	1.44	1.83	2.83				
	HEMBA1005732	4.64	3.73	2.82	4.17	4.78	5.5	3.41	2.84	3.27				
	HEMBA1005737	2.11	1.17	0.89	1.64	1.86	1.55	2.37	1.99	1.73				
	HEMBA1005742	2.91	1.85	1.65	20.12	22.7	20.93	10.11	6.75	7.19	**	+	**	+
45	HEMBA1005746	3.55	2.22	2.55	2.88	5.21	3.91	2.28	2.67	1.49				
	HEMBA1005747	6.73	2.98	3.61	4.2	6.34	4.06	4.88	4.78	5.21				
	HEMBA1005749	16	15.05	7.61	16.72	17.56	14.78	13.73	10.17	19.02				
	HEMBA1005755	1.55	1.38	0.58	2.76	3.45	1.74	2.11	2.82	2.29		*	+	
	HEMBA1005760	6.22	4.23	3.01	5.27	5.19	5.24	4.36	3.24	4.73				
50	HEMBA1005765	5.47	4.02	4.47	8.82	8.58	6.98	4.72	5.79	3.58	**	+		
	HEMBA1005766	6.49	3.72	3.07	6.86	5.34	6.17	4.5	5.2	3.85				
	HEMBA1005780	5.24	3.72	3.56	7.77	10.48	12.03	5.65	6.93	5.8	*	+	*	+
	HEMBA1005795	2.44	2.1	2.01	3.69	3.63	2.88	1.69	3.18	1.68	*	+		
	HEMBA1005809	23.36	22	11.6	14.58	20.18	18.5	16.89	18.97	9.81				
	HEMBA1005813	3.44	3.32	2.49	3.52	4.47	4.04	2.83	4.45	3.63				
55	HEMBA1005815	6.13	3.52	2.7	5.29	7.35	4.96	4.74	5.46	7.01				
	HEMBA1005822	4.2	1.96	2.92	8.67	7.02	9.4	4.99	3.69	6.16	**	+		

Table 194

	HEMBA1005317	1.33	0.36	0.19	4.23	3.8	4.6	1.25	1.46	1.92	**	+		
	HEMBA1005318	1.08	0.85	0.59	0.97	1.89	1.29	1.43	1.82	1.13				
5	HEMBA1005324	3.04	2.4	1.83	6.59	7.62	7.75	5.26	6.51	7.55	**	+	**	+
	HEMBA1005331	0.95	1.56	1.2	1.7	1.65	2.13	0.66	2.53	0.91				
	HEMBA1005337	2.8	1.37	1.32	2.67	3.1	2.37	2.01	2.34	2.18				
	HEMBA1005338	4.38	1.6	2.45	4.11	1.92	3.95	3.55	3.33	3.26				
	HEMBA1005344	22.24	11.71	11.54	14.09	14.09	14.6	12.65	14.29	14.22				
10	HEMBA1005353	6.55	4.18	3.72	6.77	13.54	9.81	6.95	6.75	7.1				
	HEMBA1005359	7.54	5.12	6.63	11.85	12.2	12.76	7.38	8.41	9.39	**	+		
	HEMBA1005362	9.18	7.14	7.14	5.77	8.95	8.4	3.09	3.31	2.6			**	-
	HEMBA1005364	0.89	1.26	0.41	1.96	2.44	1.02	1.19	1.6	1.68				
	HEMBA1005367	3.22	2.29	1.05	4.88	6.98	6.68	5.63	8	6.43	*	+	**	+
15	HEMBA1005372	2.2	0.98	0.77	1.74	3.83	3.08	4.16	2.78	2.66			*	+
	HEMBA1005374	6.99	3.71	3.35	12.54	10.52	8.75	6.1	6.58	7.22	*	+		
	HEMBA1005379	1.84	1.63	1.2	1.2	1.49	2.65	1.75	1.09	1.97				
	HEMBA1005382	7.86	4.67	5.2	10.89	7.83	8.14	5.58	6.98	6.52				
	HEMBA1005384	4.42	2.21	2.13	6.74	6.14	5.84	4.87	4.21	4.01	*	+		
20	HEMBA1005386	6.04	3.65	3.38	6.45	5.92	6.1	5.2	4.67	5.78				
	HEMBA1005389	5.36	3.94	2.77	5.75	6.88	6.02	2.6	5.56	3.66				
	HEMBA1005394	6.27	3.67	3.58	3.93	4.59	4.22	2.21	4.81	3.15				
	HEMBA1005403	11.32	8.45	6.9	16.3	23.03	11.57	16.03	13.06	13.2			*	+
	HEMBA1005408	4.6	4.51	2.17	5.61	4.87	4.3	5.51	3.2	4.27				
25	HEMBA1005410	1.48	1.46	0.98	2.22	1.83	2.32	3.82	2.31	2.31	*	+	*	+
	HEMBA1005411	3.32	2.25	1.72	8.56	7.19	8.45	4.84	3.85	4.74	**	+	*	+
	HEMBA1005423	4.84	2.65	2.83	7.04	5.69	5.75	3.26	4.32	3.64	*	+		
	HEMBA1005426	1.66	0.94	1.03	2.84	2.24	2.73	1.74	2.79	1.34	**	+		
	HEMBA1005427	18.06	13.04	14.1	24.89	25.18	27.94	11.55	18.31	15.99	**	+		
30	HEMBA1005430	3.16	1.5	2.13	1.75	2.9	3.37	2.43	3.98	2.23				
	HEMBA1005438	4.91	3.54	3.44	5.97	8.41	5.02	5.97	4.67	6.58				
	HEMBA1005443	11.24	11.79	6.21	19.21	19.58	15.66	17.03	13.17	10.83	*	+		
	HEMBA1005447	3.13	3.2	1.74	4.18	4.12	4.68	2.92	2.36	2.86	*	+		
	HEMBA1005449	4.87	2.92	3.15	2.75	4.63	3.51	2.81	3.38	5.99				
35	HEMBA1005452	8.28	4.39	4.04	3.56	7.29	6.13	4.29	5.16	4.62				
	HEMBA1005454	6.03	4.13	3.77	3.63	4.31	5.36	2.84	5.74	3.18				
	HEMBA1005468	8.63	4.08	5.4	8.19	9.91	9.17	5.46	7.18	6.46				
	HEMBA1005469	7.04	4.49	4.09	8.04	6.87	9.35	3.55	5.47	4.98				
	HEMBA1005472	4.58	4.13	2.33	5.09	7.14	6.31	4.57	3.72	5.09				
40	HEMBA1005474	7.99	6.35	8.53	12.45	17.71	14.57	6.84	6.03	7.86	*	+		
	HEMBA1005475	27.06	16.75	12.04	21.27	20.2	24.59	14.7	11.72	14.55				
	HEMBA1005489	4.67	3.91	3.31	12.33	12.95	12.78	5.02	3.73	4.43	**	+		
	HEMBA1005497	1.7	0.87	0.7	1.28	2.32	1.65	1.49	1.73	0.9				
45	HEMBA1005500	6.11	2.66	2.28	6.01	8.49	7.76	2.99	5.44	4.21				
	HEMBA1005506	1.91	0.96	0.87	1.02	1.78	1.61	1.14	3.14	1.21				
	HEMBA1005508	3	1.68	2.62	3.65	3.78	4.7	1.31	2.01	2	*	+		
	HEMBA1005511	6.78	4.02	3.71	12.46	10.15	10.8	6.67	5.32	7.37	**	+		
	HEMBA1005513	9.39	4.07	4.88	7.16	6.69	8.41	5.04	6.55	4.92				
	HEMBA1005517	4.77	2.9	3.52	2.59	3.48	4.27	1.92	3.32	2.46				
	HEMBA1005518	6.02	2.95	2.57	4.55	4.62	5.87	5.99	3.9	5.53				
50	HEMBA1005520	11.23	5.82	6.06	14.5	18.42	18.84	7.99	9.11	9.67	*	+		
	HEMBA1005522	4.58	1.74	1.96	2.63	3.4	3.05	1.78	3.18	2.26				
	HEMBA1005526	4	2.06	4.25	8.46	10.15	10.58	3.26	5.34	5.19	**	+		
	HEMBA1005528	14.83	10.54	9.95	13.03	18.88	16.21	6.65	7.84	6.97			*	-
	HEMBA1005530	5.44	2.29	3.17	4.84	6.25	8.18	4.67	4.21	3.84				
55	HEMBA1005538	4.71	2.93	2.46	83.2	102.3	97.16	227.3	162.3	210.6	**	+	**	+
	HEMBA1005539	7.02	4.61	3.84	4.34	5.62	5.7	5.14	4.99	5.58				
	HEMBA1005545	4.05	4.59	3.18	3.31	5.22	4.49	4.33	4.46	3.97				

Table 193

	HEMBA1005050	8.01	4.69	4.35	6.4	8.24	6.75	4.64	5.95	4.47				
	HEMBA1005062	2.24	3.49	0.58	2.31	2.34	1.56	1.28	2.55	1.35				
5	HEMBA1005066	1.59	0.53	1.22	1.43	2.19	2.08	0.94	1.37	2.12				
	HEMBA1005067	10.97	5.24	5.8	11.93	6.24	15.81	3.48	6.87	4.63				
	HEMBA1005070	54.34	32.66	23.12	7.48	7.23	9.46	4.96	6.22	5.67	*	-	*	-
	HEMBA1005075	4.78	2.93	2.39	9.53	8.99	8.84	5.77	6.79	5.19	**	+	*	+
	HEMBA1005078	9.58	7.81	5.77	9.39	9.72	10.05	5.01	5.86	6.72				
10	HEMBA1005079	12.04	7.57	6.48	19.42	17.72	15.43	8.5	9.75	11.45	*	+		
	HEMBA1005083	2.66	1.46	0.66	1.94	3.02	2.07	1.27	2.07	1.73				
	HEMBA1005084	7.91	6.72	4.77	5.71	7.85	8.74	5.49	4.34	5.97				
	HEMBA1005088	2.86	1.68	1.86	2.41	5.46	5.18	1.67	3.51	1.49				
	HEMBA1005089	5.98	4.14	4.5	9.36	10.56	9.53	2.93	5.59	4.57	**	+		
15	HEMBA1005090	33.54	22.43	17.55	44.06	43.43	42.47	20.48	22.61	17.3	*	+		
	HEMBA1005096	5.76	3.96	4.37	6.03	5.87	6.22	4.23	3.3	6.27				
	HEMBA1005101	5.71	2.76	3.85	3.75	5.23	3.72	2.48	3.36	3.77				
	HEMBA1005107	4.5	1.82	2.91	2.69	3.89	3.12	3.06	3.74	2.52				
	HEMBA1005113	1.43	0.81	0.45	8.23	11.09	10.71	5.43	5.23	6.51	**	+	**	+
20	HEMBA1005123	10.61	5.86	5.3	15.09	21.59	18.64	8.57	8.85	8.33	*	+		
	HEMBA1005133	2.6	2.55	2.08	5.44	6.93	6.67	3.17	4.12	2.67	**	+		
	HEMBA1005135	1.91	1.13	1.66	1.75	3.38	1.54	1.31	3.02	1.14				
	HEMBA1005145	16.67	9.87	9.21	12.39	15.8	16.28	8.2	10.27	10.61				
	HEMBA1005149	10.32	5.61	5.06	11.44	12.5	12.06	7.17	8.59	7.71	*	+		
25	HEMBA1005152	6.34	4.06	3.55	9.52	11.4	12.18	3.28	4.04	5.79	**	+		
	HEMBA1005159	0.7	1.49	0.94	1.57	2.36	1.76	1.22	2.54	0.91				
	HEMBA1005172	43.22	25.23	24.37	33.5	39.86	37.96	32.09	25.74	34.44				
	HEMBA1005185	4.97	4.57	2.99	2.86	3.27	4.08	2.48	3.14	1.7				
	HEMBA1005186	3.35	2.42	3.23	5.64	6.25	4.46	2.06	2.21	2.79	*	+		
	HEMBA1005195	1.99	0.84	0.81	1.89	2.31	1.52	1.31	2.87	1.25				
30	HEMBA1005201	6.2	5.19	2.55	5.77	6.88	6.27	4.89	5.22	6.55				
	HEMBA1005202	8.96	4.63	5.23	6.96	8.01	6.67	8.1	7.62	9.46				
	HEMBA1005204	113.3	93.42	81.36	145.9	165	106.5	90.09	59.5	89.11				
	HEMBA1005206	6.48	3.93	4.87	5.9	5.71	6.15	4.98	4.32	4.52				
	HEMBA1005219	2.14	1.72	1.8	4.03	2.98	2.85	3.28	4.04	4.31	*	+	**	+
35	HEMBA1005223	3.02	2.16	2.78	4.29	3.41	4.21	2.9	3.66	3.28	*	+		
	HEMBA1005229	0.71	0.07	0.59	1.25	1.02	0.47	0.51	2.08	0.98				
	HEMBA1005230	4.24	4.62	2.37	7.34	7.76	6.64	2.52	4.81	4.22	*	+		
	HEMBA1005232	0.15	0.54	0.47	1.05	1.44	1.37	1.1	0.73	0.86	**	+	*	+
	HEMBA1005238	5.05	3.37	2.42	6.46	5.11	6.11	4.05	3.86	3.91				
40	HEMBA1005241	18.2	11.3	9.41	11.74	14.66	18	9.85	7.33	9.11				
	HEMBA1005244	6.45	3.35	4.4	5.3	7.24	5.85	3.98	5	6.42				
	HEMBA1005246	9.39	6.95	6.65	15.52	17.83	13.37	15.28	9.28	12.96	**	+		
	HEMBA1005251	2.49	1.43	2.18	5.25	6.15	4.92	3.41	3.93	3.35	**	+	*	+
	HEMBA1005252	3.83	2.63	3.03	3.56	4.92	3.46	2.88	4.5	4.38				
45	HEMBA1005267	1.63	0.84	1.67	10.27	7.55	7.28	1.17	3.13	1.81	**	+		
	HEMBA1005274	1.18	0.71	0.61	1.46	2.14	1.62	1.18	1.08	1.02	*	+		
	HEMBA1005275	1.9	0.81	0.85	2.82	4.11	3.4	1.92	2.54	1.27	*	+		
	HEMBA1005288	3.5	1.84	2.36	6.6	8.72	6.93	3.45	3.43	3.32	**	+		
	HEMBA1005293	1.91	2.03	0.55	4.54	1.95	1.33	0.58	2.15	1.2				
50	HEMBA1005296	401.9	314.1	296.6	377.8	403.7	432.4	228.1	207.4	230		*	-	
	HEMBA1005301	1.98	0.74	1.57	2.67	1.35	1.8	1.62	2.6	1.33				
	HEMBA1005304	4.1	2.37	2.93	8.99	8.69	9.63	4.7	6.26	5.96	**	+	*	+
	HEMBA1005305	2.8	1.25	1.81	4.03	4.81	4.66	2.57	4.35	2.78	**	+		
	HEMBA1005311	2.04	1.03	1.55	2.81	2.74	3.45	0.88	2.3	2.36	*	+		
	HEMBA1005313	6.91	3.99	3.19	6.31	4.42	4.78	4.14	4.74	6.56				
55	HEMBA1005314	0.55	0.27	0.2	1.02	1.14	0.89	1.2	0.4	1.03	**	+		
	HEMBA1005315	4.12	1.27	1.36	3	4.13	3.44	3.48	2.53	3.28				

Table 192

	HEMBA1004778	4.28	3.09	3.12	5.87	7.81	8.46	5.37	4.86	3.66	*	+		
	HEMBA1004784	1.55	1.14	0.87	1.97	2.67	2.4	1.81	2.87	1.66	*	+		
5	HEMBA1004785	2.2	0.85	1.41	2.94	2.11	2.82	2.94	3.76	2.42			*	+
	HEMBA1004789	2.02	2.15	2.94	6	4.66	4.07	4.12	6.23	6.59	*	+	*	+
	HEMBA1004795	1.94	0.91	1.99	4.74	2.62	2.39	1.99	2.85	2.46				
	HEMBA1004797	3.34	1.51	1.57	3.19	4.14	4.19	3.42	2.94	4.14				
10	HEMBA1004803	1.73	1.53	0.52	3.19	3.28	3.24	3.3	2.11	2.68	**	+	*	+
	HEMBA1004806	1.99	0.24	0.76	2.51	2.13	1.62	1.14	2.33	1.41				
	HEMBA1004807	6.07	4.25	4.5	4.85	8.03	9.33	4.48	5.59	5.41				
	HEMBA1004816	3.49	2.36	1.89	3.34	3.8	3.31	2.37	4.02	1.69				
	HEMBA1004820	1.49	1.14	1.32	2.51	2.88	2.8	1.5	4.47	1.86	**	+		
15	HEMBA1004833	7.98	3.57	4.1	7.09	8.03	7.72	4.99	7.63	6.59				
	HEMBA1004847	6.33	4.11	5.21	8.38	7.16	8.48	4.35	8.93	6.34	*	+		
	HEMBA1004850	3.92	2.57	2.41	5.26	3.09	3.63	3.54	3.4	5.81				
	HEMBA1004863	4.26	1.79	2.07	6.34	5.16	5.37	2.36	2.91	5.42	*	+		
	HEMBA1004864	8.29	3.32	3.08	5.48	7.27	7.94	4.75	3.71	4.59				
20	HEMBA1004865	1.92	1.18	0.62	2.11	6.7	3.86	2.14	1.94	1.68				
	HEMBA1004880	4.54	3.09	3.36	6.03	7.12	7.25	3.5	4.7	4.49	**	+		
	HEMBA1004882	5.35	4.05	3.06	4.2	4.72	3.45	2.62	4.51	3.09				
	HEMBA1004885	1.17	0.68	0.57	1.14	0.82	0.86	0.53	1	0.47				
	HEMBA1004889	3.26	2.08	1.7	3.09	2.94	3.37	2.23	2.83	5.72				
25	HEMBA1004900	1.39	1.1	0.25	1.7	1.35	1.1	1.57	1.47	1.61				
	HEMBA1004909	6.14	4.05	3.74	6.91	8	7.96	4.94	4.32	5.82	*	+		
	HEMBA1004918	4.98	2.15	2.73	5.38	6.39	6.51	3.65	3	3.79	*	+		
	HEMBA1004923	1.88	1.64	1.69	3.18	2.96	3.02	2.23	2.61	2.53	**	+	**	+
	HEMBA1004929	2.42	1.04	1.11	2.68	2.08	2.3	2.43	1.05	1.27				
30	HEMBA1004930	5.54	5.02	5.16	8.04	11.27	11.38	5.24	6.2	5.58	*	+		
	HEMBA1004933	2.24	1.54	1.06	2	2.4	2.08	1.19	1.47	2.06				
	HEMBA1004934	0.55	0.77	0.07	1.15	0.99	1.58	1.85	2.74	1.58	*	+	*	+
	HEMBA1004937	6.5	2.53	3.22	3.69	3.97	5.19	4.16	4.2	3.69				
	HEMBA1004943	6.44	2.93	2.55	5.45	3.9	5.9	3.81	4.39	5.14				
35	HEMBA1004944	4.47	1.97	2.6	5.4	4.69	6.01	3.98	3.08	5.3	*	+		
	HEMBA1004946	6.58	4.26	2.56	8.23	7.78	9.16	5.73	6.06	6.35	*	+		
	HEMBA1004952	5.05	2.8	1.43	3.17	3.75	3.4	2.89	3.56	3.26				
	HEMBA1004954	2.94	2.13	2.53	7.6	9.09	8.39	8.28	11.47	6.83	**	+	*	+
	HEMBA1004956	1.7	0.98	0.85	2.16	2.35	1.65	2.19	1.65	0.68				
	HEMBA1004960	4.22	1.35	1.83	3.33	4.35	3.89	3.18	2.33	2.62				
40	HEMBA1004971	2.85	2.08	2.33	3.11	3.19	2.48	4.48	3.31	3.12				
	HEMBA1004972	7.97	3.44	5.28	7.05	7.91	7.94	4.91	4.41	4.71				
	HEMBA1004973	4.05	2.96	1.6	4.3	3.46	4.03	3.1	2.58	3.76				
	HEMBA1004977	14.24	10.04	6.48	10.74	14.23	17.72	5.8	5.62	5.43				
	HEMBA1004978	3.63	3.21	1.82	4.34	4.05	5.53	3.79	4.18	2.53				
45	HEMBA1004980	2.51	2.43	1.78	4.29	5.14	5.81	2.73	3.03	2.97	**	+		
	HEMBA1004982	1.4	0.95	0.55	1.1	1.4	2.11	0.94	2.12	0.81				
	HEMBA1004983	1.7	1.5	1.07	1.2	1.85	1.29	1.7	1.13	1.37				
	HEMBA1004995	4.75	4.53	4.44	5.51	5.64	4.6	3.99	3.69	4.53				
	HEMBA1005004	4.11	3.34	2.48	4.8	3.41	3.91	3.87	2.23	2.59				
	HEMBA1005008	5.55	2.4	3.38	3.53	5.55	4.97	3.01	4.08	3.22				
50	HEMBA1005009	10.15	9.95	7.66	7.94	11.36	8.06	4.21	5.06	5.68		**	-	
	HEMBA1005019	6.33	2.93	3.49	4.34	5.65	6.43	4.58	6.41	4.98				
	HEMBA1005021	5.34	2.42	3.36	5.76	5.02	6.07	3.05	4.52	3.35				
	HEMBA1005029	7.09	2.85	5.15	5.79	6.77	7.31	4.12	4.52	4.02				
	HEMBA1005035	13.39	11.61	9.27	16.88	22.86	20.98	13.14	12.08	14.81	*	+		
55	HEMBA1005036	9.37	4.9	6.57	4.71	7.3	8.39	7.97	8.49	9.16				
	HEMBA1005039	2.56	2.26	1.97	3.46	4.91	5.18	2.9	3.33	3.77	*	+	*	+
	HEMBA1005047	3.73	2.69	2.58	2.7	3.22	4.69	3.19	3.28	3.52				

Table 191

	HEMBA1004566	28.53	23.96	23.72	13.72	19.42	18.07	12.15	16.7	14.58	*	-	**	-
	HEMBA1004573	2.19	1.72	1.51	3.93	5.22	5.71	3.32	3.47	1.99	**	+		
5	HEMBA1004576	2.94	1.45	1.92	18.03	33.01	34.57	7.81	8.41	10.12	**	+	**	+
	HEMBA1004577	5	2.83	2.54	7.07	10.98	8.6	4.13	8.82	4.99	*	+		
	HEMBA1004586	5.72	3.41	4.19	8.7	7.46	11.19	4.1	6.11	4.48	*	+		
	HEMBA1004596	4.81	2.28	2.02	2.98	3.67	3.46	2.47	2.61	3				
	HEMBA1004604	6.48	4.01	3.96	4.74	6.55	5.9	8.49	6.15	4.83				
10	HEMBA1004607	3.7	2.23	1.35	4.64	5.86	4.48	2.81	3.79	3.93	*	+		
	HEMBA1004610	4.03	2.57	2.33	4.52	5.94	4.83	3.02	2.81	3	*	+		
	HEMBA1004617	2.21	4.92	1	2.84	9.03	3.69	1.85	2.86	2.94				
	HEMBA1004622	5.45	3.28	2.52	5.48	8.39	9.1	4.14	4.48	4.68				
	HEMBA1004626	4.11	2.56	2.25	5.1	4.71	5.91	2.73	4.36	3.32	*	+		
15	HEMBA1004629	3.07	1.77	1.42	3.68	3.77	4.82	1.19	3.75	1.18	*	+		
	HEMBA1004631	1.43	2.39	0.95	2.12	1.94	2.84	2.88	1.6	2.44				
	HEMBA1004632	2.27	1.83	1.79	2.78	2.92	1.76	2.34	3.5	2				
	HEMBA1004633	7.83	5.66	4.81	4.47	6.1	5.15	5.55	4.15	5.55				
	HEMBA1004636	6.11	4.03	3.37	5.56	5.52	5.5	4.94	4.1	4.16				
20	HEMBA1004637	3.8	2.43	1.85	2.17	3.96	3.28	2.95	2.5	2				
	HEMBA1004638	1.58	0.7	0.19	0.85	2.26	3.04	1.06	1.19	1.64				
	HEMBA1004645	4.58	1.72	2.46	3.58	5.23	5.82	2.85	4.55	3.78				
	HEMBA1004656	3.49	2.49	3.49	3.55	3.42	3.65	2.19	3.03	2				
	HEMBA1004657	23.62	14.49	14.4	48.51	47.67	43.85	51.21	56.08	58.34	**	+	**	+
25	HEMBA1004666	1.8	1.42	1.03	2.78	2.47	2.72	1.97	2.35	2.06	**	+	*	+
	HEMBA1004669	5.4	3.16	2.59	6.16	6.23	6.59	2.94	2.65	2.66	*	+		
	HEMBA1004670	4.37	2.24	2	5.27	6.01	4.17	2.94	3.39	4.41				
	HEMBA1004672	5.55	2.84	2.98	5.68	8.28	8.14	3.49	6.01	3.36	*	+		
	HEMBA1004689	43.34	14.93	30.58	21.98	24.65	26.05	13.05	14.68	11.72				
	HEMBA1004690	4.61	2.61	2.69	2.94	2.18	2.84	1.97	4.01	2.41				
30	HEMBA1004693	2.15	1.25	1.33	2.01	3.2	3.06	1.39	3.08	2				
	HEMBA1004697	7.39	3.61	2.79	5.75	7.36	9.2	5.36	4.7	6.2				
	HEMBA1004702	21.02	14.02	11.62	9.2	10.6	12.82	11.9	12.65	12.79				
	HEMBA1004704	6.08	3.81	3.24	8.5	8.45	8.19	4.75	5.52	5.39	*	+		
	HEMBA1004705	1.15	0.61	0.21	1.49	1.26	1.73	1.37	1.44	1.36				
35	HEMBA1004706	3.9	2.72	2.07	2.01	3.27	2.47	3.18	2.94	2.37				
	HEMBA1004709	3.4	2.4	2.61	5.18	5.97	7.11	2.19	3.92	3.09	**	+		
	HEMBA1004711	3.02	1.29	2.07	2.19	3.65	3.64	1.38	3.22	1.38				
	HEMBA1004723	9.52	5.41	7.44	9.15	11.88	10.6	5.92	9.59	6.27				
	HEMBA1004725	5.24	3.87	3.31	6.21	5.61	5.19	5.65	5.85	6.52		*	+	
40	HEMBA1004730	1.7	2.99	1.13	11.04	3.71	3.48	1.14	4.24	1.15				
	HEMBA1004733	1.86	1.11	1.27	1.93	2.88	2.54	1.38	2.89	2.03	*	+		
	HEMBA1004734	2.06	1.99	1.5	2.15	2.83	2.85	2.1	2.82	2.29				
	HEMBA1004736	3.46	3.3	2.73	5.69	8.26	7.15	2.94	3.83	4.08	**	+		
	HEMBA1004748	4.24	1.57	1.93	4.83	6.28	6.83	2.64	4.21	2.49	*	+		
45	HEMBA1004749	7.35	4.59	5.33	5.23	6.38	10.26	4.24	7.9	6.41				
	HEMBA1004751	3.74	2.05	2.99	5.29	6.07	7.15	2.9	5.44	3.62	*	+		
	HEMBA1004752	5.63	3.05	2.11	4.83	5.66	7.24	4.55	3.43	5.1				
	HEMBA1004753	85.27	60.35	45.13	73.61	76.67	82.04	35.88	33.51	34.18				
	HEMBA1004755	12.21	10.42	8.56	18.13	22.58	19.53	19.43	13.65	17	**	+	*	+
50	HEMBA1004756	1.98	0.4	0.9	1.17	2.37	1.88	1.4	2.86	2.34				
	HEMBA1004758	3.05	2.33	2.23	5.05	4.15	4.14	2.36	3.68	2.72	**	+		
	HEMBA1004763	2.53	2.54	2.42	3.64	3.57	2.87	2.52	4.35	2.6	*	+		
	HEMBA1004768	0.63	0.57	0.48	2.03	2.91	1.86	1.11	2.85	0.94	**	+		
	HEMBA1004770	1.17	0.28	1.04	3.43	2.96	3.94	2.26	2.05	1.62	**	+	*	+
	HEMBA1004771	3.01	1.5	1.36	3.25	3.04	3.46	2.2	2.12	2.05				
55	HEMBA1004775	6.8	4.62	3.7	7.13	8.07	9.04	7.62	7.16	8.82	*	+		
	HEMBA1004776	3.71	2.57	1.18	9.61	3.42	2.31	3.1	3.48	4.38				

Table 190

	HEMBA1004329	6.64	4.05	3.69	10.1	11.36	10.31	6.59	6.39	7.21	**	+		
5	HEMBA1004330	3.08	1.92	1.4	2.71	3.14	3.34	2.64	3.02	3.26				
	HEMBA1004334	3.9	1.95	1.91	3.91	3.51	4.57	2.69	2.68	1.57				
	HEMBA1004335	4.91	2.24	2.81	7.41	9.8	7.49	5.19	6.78	5.7	*	+		
	HEMBA1004341	6.84	4.27	5.5	4.53	6.2	5.05	5.02	6.83	6.93				
	HEMBA1004344	17.75	13.13	14.74	15.95	19.67	19.67	18.23	19.51	24.66				
10	HEMBA1004347	4.63	3.35	2.01	5.16	6.48	5.36	2.73	3.19	3.71				
	HEMBA1004349	8.89	2.46	3.99	12.23	16.69	10.37	7.71	6.98	8.61	*	+		
	HEMBA1004352	5.41	3.1	3.3	7.91	8.12	10.45	4.93	5.52	5.36	*	+		
	HEMBA1004353	8.35	7.6	6.31	15.51	16.38	15.96	6.75	8.35	9.31	**	+		
	HEMBA1004354	4.38	1.54	2.32	5.25	5.81	6.37	3.27	4.92	3.61	*	+		
15	HEMBA1004356	2.81	2.85	3.03	5.06	4.66	5.46	5.28	5.77	4.17	**	+	*	+
	HEMBA1004360	5.79	2.16	5.01	6.93	5.95	5.72	3.15	6.55	5.08				
	HEMBA1004366	2.78	2.3	2.86	5.4	6.73	4.61	2.18	3.01	3.38	*	+		
	HEMBA1004372	0.38	0.27	0.43	0.47	0.63	0.99	0.52	0.83	0.34				
	HEMBA1004377	7.38	3.14	3.85	11.65	12.1	15.48	9.22	8.78	11.95	**	+	*	+
20	HEMBA1004389	18.67	11.71	10.38	8.69	8.39	17.15	9.23	8.15	7.38				
	HEMBA1004391	2.93	2.48	2.45	7.42	5.09	7.12	3.62	4.64	3.41	**	+	*	+
	HEMBA1004393	18.44	14.15	13.12	19.38	17.77	18.16	22.31	14.59	20.28				
	HEMBA1004394	1.18	1.11	1.72	2.3	1.6	2.38	1.09	4.42	1.46				
	HEMBA1004396	1.79	1.02	1.22	3.41	3.48	3.73	1.3	3.02	1.73	**	+		
25	HEMBA1004401	4.73	3.38	4.96	4.16	4.54	5.13	2.63	5.44	3.27				
	HEMBA1004405	3.95	2.13	1.81	6.15	8.26	6.59	3.78	4.33	5.63	**	+		
	HEMBA1004408	5.72	3.65	3.17	5.44	6.45	4.46	2.34	3.68	2.97				
	HEMBA1004414	8.38	4.86	5.28	9.94	19.52	21.58	6.98	7.48	7.61	*	+		
	HEMBA1004429	3.38	2.07	1.78	8.58	8.61	9.23	4.27	3.18	4.51	**	+		
	HEMBA1004433	1.82	1.56	1.04	5.34	5.56	5.46	1.92	2.85	2.38	**	+		
30	HEMBA1004440	2.19	0.58	1.67	2.76	2.16	2.15	1.08	2.89	1.62				
	HEMBA1004444	4.28	2	2.33	6.71	7.29	10.11	5.5	5.93	3.39	*	+		
	HEMBA1004446	1.19	0.41	1.18	2.01	2.51	2.6	0.58	1.63	1.83	*	+		
	HEMBA1004451	4.92	5.14	2.78	5.62	4.16	5.1	2.95	3.75	4.07				
	HEMBA1004452	1.45	1.3	0.96	7.34	8.28	11.36	3.26	5.07	5.69	**	+	**	+
35	HEMBA1004454	2.75	3.17	2.58	3.68	3.73	5.7	3.62	3.63	3.66		*	+	
	HEMBA1004460	8.77	5.29	4.63	9.49	11.6	11.51	5.17	5.78	6.91	*	+		
	HEMBA1004461	3.02	1.29	1.56	1.22	2.06	2.62	1.48	2	2.51				
	HEMBA1004468	9.69	5.12	5.83	5.76	9.08	12.25	6.18	7.22	5.91				
	HEMBA1004479	5.17	2.6	2.53	3.06	4.8	5.24	1.98	4.08	3.44				
	HEMBA1004482	2.81	3.98	3.7	2.47	3.92	2.52	2.59	2.29	3.11				
40	HEMBA1004491	1.37	1	0.96	1.25	1.97	1.96	0.89	1.47	2.84				
	HEMBA1004499	6.22	5.75	3.57	9.95	9.17	8.62	6.22	6.62	6.45	*	+		
	HEMBA1004502	3.1	2.59	1.77	4.11	5.34	4.51	4.17	2.98	4.03	*	+		
	HEMBA1004505	4.8	2.59	1.93	2.42	4.25	3.38	2.91	2.43	2.1				
45	HEMBA1004506	2.39	1.28	1.21	2.96	3.46	3.27	2.23	2.51	1.92	*	+		
	HEMBA1004507	70.44	39.05	46.26	43.39	51.75	50.62	19.17	24.55	22.93		*	-	
	HEMBA1004509	5.46	3.62	4.71	3.53	4.82	5.37	2.96	3.83	2.3				
	HEMBA1004523	1.41	0.75	0.59	1.16	1.53	1.37	1.32	1.24	1.24				
	HEMBA1004528	3.19	1.97	1.1	3.38	4.01	3.33	4.31	3.09	4.88				
	HEMBA1004534	6.12	2.73	4	6.77	8.18	7.93	6.04	5.56	6.21	*	+		
50	HEMBA1004536	4.76	3.38	3.05	3.55	4.6	4.52	2.5	2.23	2.99				
	HEMBA1004538	21.21	15.5	13.77	31.9	33.44	32.76	19.4	20.15	17.02	**	+		
	HEMBA1004542	2.99	2.19	1.59	3.03	3.58	3.02	3.51	3.43	2.25				
	HEMBA1004552	7.56	6.12	5.53	7.59	13.46	14.87	4.88	6.44	7.28				
	HEMBA1004554	2.07	2.28	0.95	2.8	2.16	2.43	2.95	3.09	3.26		*	+	
55	HEMBA1004558	11.57	6.62	6.21	7.21	8.48	8.56	6.35	6.8	7.65				
	HEMBA1004560	4.78	3.27	2.78	3.55	5.31	4.2	5.01	4.88	3.58				
	HEMBA1004564	7.43	4.79	5.05	12.74	14.02	11.12	5.94	7.39	6.24	**	+		

Table 189

	HEMBA1004110	14.95	6.9	7.32	18.8	19.15	18.28	8.68	9.12	8.42	*	+		
5	HEMBA1004111	4.86	3.05	3.79	7.76	8.14	8.2	4.79	4.58	4.21	**	+		
	HEMBA1004124	6.94	3.7	4.42	6.1	3.89	4.52	4.49	4.21	4.75				
	HEMBA1004130	9.54	3.62	3.55	9.36	10.43	9.03	4.61	5.66	5.54				
	HEMBA1004131	4.85	3.97	3.36	3.86	4.83	4.69	2.77	4.06	2.93				
	HEMBA1004132	3.06	2.1	4.1	5.2	8.61	9.33	4.36	6.28	4.36	*	+		
10	HEMBA1004133	4.53	2.37	1.71	4.62	2.8	5.4	3.39	2.94	3.1				
	HEMBA1004138	4.15	2.09	2.18	3.1	3.45	2.71	3.21	2.5	3.12				
	HEMBA1004143	5.3	2.88	3.04	7.57	5.72	6.31	5.7	5.05	4.79	*	+		
	HEMBA1004146	4.2	1.65	2.04	5.44	5.59	4.27	2.49	3.4	3.26				
	HEMBA1004148	6.71	2.61	2.68	3.24	3.23	4.49	2.38	4.66	3.54				
15	HEMBA1004149	1.73	0.7	0.91	2.13	2.29	1.85	1.73	1.38	1.4	*	+		
	HEMBA1004150	1.14	0.72	0.48	0.29	0.95	0.79	0.76	1.47	0.27				
	HEMBA1004154	10.52	5.49	6.9	6.41	6.4	7.33	4.84	8.66	5.4				
	HEMBA1004164	7.02	3.4	3.27	9.28	10.11	8.81	5.16	5.49	5.3	*	+		
	HEMBA1004168	11.84	7.61	6.03	3.62	4.76	4.13	3.34	2.92	1.89		*	-	
20	HEMBA1004199	0.92	0.62	0.74	1.67	1.3	1.87	1.47	1.95	1.4	*	+	*	+
	HEMBA1004200	1.57	1.23	0.4	3.73	3.83	2.53	1.99	2.05	1.43	*	+		
	HEMBA1004201	3.89	3.07	2.03	3.6	4.1	3.75	2.3	4.41	3.45				
	HEMBA1004202	4.9	3.79	2.27	2.88	2.3	2.76	2.58	3.69	2.7				
	HEMBA1004203	5.77	1.33	2.87	2.88	5.01	4.03	2.05	3.68	2.38				
25	HEMBA1004207	0.56	0.3	0.47	1.04	1.74	1.87	0.89	2.13	1.31	*	+		
	HEMBA1004210	8.61	6.61	6.14	2.77	2.66	3.95	1.89	2.45	2.61	**	-	**	-
	HEMBA1004225	5.03	3.74	3.98	8.75	9.84	8.6	5.26	5.3	4.25	**	+		
	HEMBA1004227	3.79	2.62	4.1	4.2	4.62	3.31	3.12	4.12	2.59				
	HEMBA1004235	7.02	4	4.13	5.38	8.95	5.38	4.38	6.22	4.55				
	HEMBA1004237	3.9	2.42	2.47	3.59	4.2	5.86	2.12	3.58	2.89				
30	HEMBA1004238	6.25	1.89	3.24	4.96	7.33	6.03	3.76	4.17	3.98				
	HEMBA1004241	0.67	0.27	0.46	0.34	1.31	1.04	0.22	1.55	0.61				
	HEMBA1004242	32.46	19.09	20.5	23.42	40.5	41.44	12.31	21.44	17.84				
	HEMBA1004243	13.89	7.41	6.2	5.78	8.65	6.42	6.33	5.94	4.6				
	HEMBA1004246	2.25	1.26	2.23	4.03	4.82	3.81	2.36	4.78	2.1	**	+		
35	HEMBA1004247	5.45	2.79	1.32	2	4.11	3.23	3.04	3.5	3.55				
	HEMBA1004248	1.69	0.88	1.09	3.22	4.63	3.53	2.79	3	3.44	**	+	**	+
	HEMBA1004250	2.2	1.27	1.09	2.31	1.66	2.1	1.9	1.53	1.16				
	HEMBA1004252	3.18	2.82	2.3	4.58	5.09	4.33	3.04	3.66	2.93	**	+		
	HEMBA1004260	6.17	5.02	5.43	14.46	16.02	13.28	2.04	6.53	5.94	**	+		
40	HEMBA1004264	2.63	0.93	1.56	1.92	3.23	2.09	0.78	1.85	0.77				
	HEMBA1004267	17.36	9.92	10.53	28.33	30.44	23.65	13.63	14.33	15.75	**	+		
	HEMBA1004272	3.25	1.51	1.9	3.88	2.89	3.11	2.45	3.01	1.7				
	HEMBA1004274	4.01	2.2	1.91	2.76	5.04	4.3	3.12	2.65	2.58				
	HEMBA1004275	7.65	2.23	3.79	6.73	5.64	5.93	3.97	4.61	4.77				
	HEMBA1004276	2.41	0.9	1.94	2.49	2.68	2.17	2.68	2.47	2.33				
45	HEMBA1004279	3.98	2.11	3.24	4.12	3.59	4.41	2.04	2.95	1.7				
	HEMBA1004284	2.55	1.22	1.55	4.17	5.87	4.34	1.28	3.05	2.74	*	+		
	HEMBA1004286	2.41	1.26	2.32	1.53	2.67	2.43	1.2	3	2.02				
	HEMBA1004289	4.95	2.88	2.44	8.79	8.57	7.77	4.32	4.66	6.4	**	+		
	HEMBA1004293	20.86	17.2	15.27	23.95	23.65	21.96	12.13	13.81	16.34	*	+		
50	HEMBA1004295	3.05	1.8	2.64	2.91	2.85	3.02	1.98	3.55	3.48				
	HEMBA1004302	0.66	0.43	0.5	1.59	1.46	1.59	1.57	2.55	1.32	**	+	*	+
	HEMBA1004306	15.93	11.78	11.41	15.21	18.98	13.88	13.95	14.44	16.14				
	HEMBA1004312	2.81	2.1	2.08	6.27	6.34	5.38	1.96	3.12	2.81	**	+		
	HEMBA1004314	2.53	1.33	1.74	4.02	5.28	5.79	1.6	3.72	2.38	**	+		
55	HEMBA1004321	6.87	2.68	4.89	6.41	11.51	10.55	3.56	5.85	5.02				
	HEMBA1004323	6.15	3.8	3.34	8.44	11.8	9.65	4.55	5.84	5.92	*	+		
	HEMBA1004327	4.25	2.43	2.21	4.5	3.91	4.05	2.91	3.47	3.95				

Table 188

	HEMBA1003904	2.78	1.45	1.43	2.77	2.17	1.59	1.47	2.66	1.69				
	HEMBA1003908	1.69	1.16	1.22	2.42	2.58	2.06	1.92	2.64	1.46	*	+		
5	HEMBA1003926	72.36	45.24	46.72	61.75	49.96	64.94	25.26	18.43	24.45			*	-
	HEMBA1003937	3.1	1.85	1.98	6.12	8.5	7.61	2.66	5.69	3.16	**	+		
	HEMBA1003939	1.28	1.62	1.87	1.85	4.47	4.22	0.72	2.97	2.45				
	HEMBA1003940	2.82	0.88	1.71	2.17	3.19	2.37	0.51	2.52	1.7				
	HEMBA1003941	4.35	2.77	1.79	1.96	4.65	3.03	2.55	3.88	2.82				
10	HEMBA1003942	2.44	1.82	1.09	3.64	3.65	2.58	2.38	2.63	2.03	*	+		
	HEMBA1003945	9.46	3.83	5.74	8.44	8.96	9.42	7.88	6.57	7.46				
	HEMBA1003949	2.14	1.99	0.59	2.89	3.58	3.78	1.92	2.25	1.36	*	+		
	HEMBA1003950	1.45	1.52	0.64	1.83	1.87	1.76	1.11	1.8	1.56				
	HEMBA1003953	1.96	0.44	1.34	3.08	3.28	3.34	1.95	3.21	1.37	*	+		
15	HEMBA1003958	6.98	4.78	4.74	10.87	13.86	10.68	4.23	6.52	6.23	**	+		
	HEMBA1003959	2.84	3.02	3.46	6.74	9.97	6.27	2.64	3.74	2.94	*	+		
	HEMBA1003960	7.33	2.27	2.98	3.59	5.1	3.92	2.8	3.92	3.79				
	HEMBA1003966	4.91	3.07	2.16	3.5	4.6	3.28	2.1	2.93	3.48				
	HEMBA1003967	5.85	3.63	2.68	3.94	3.8	3.19	1.89	3.17	2.35				
20	HEMBA1003968	3.76	2.02	2.13	4.21	6.16	3.59	4.13	4.11	3.84				
	HEMBA1003974	41.47	29.67	25.73	95.3	104.1	103.5	100	82.53	110.2	**	+	**	+
	HEMBA1003976	2.48	1.1	1.38	2.13	2.25	2.22	1.34	1.82	1.6				
	HEMBA1003977	2.19	1.38	1.4	2.42	3.02	1.57	1.86	1.96	1.93				
	HEMBA1003978	2.44	1.5	1.92	3.24	3.34	3.85	1.9	2.87	2.37	*	+		
25	HEMBA1003981	7.98	4.15	3.07	6.67	7.3	7.05	6.37	6.68	8.81				
	HEMBA1003982	6.94	4.75	3.19	18.33	22.13	23.04	19.29	21.74	19.78	**	+	**	+
	HEMBA1003985	2.27	1.26	0.95	3.01	1.91	1.85	1.02	2.35	1.03				
	HEMBA1003987	3.79	1.42	2.2	4.67	5.44	5.59	3.67	4.19	3.44	*	+		
	HEMBA1003989	2.32	1.65	1.59	4.16	4.13	5.73	3.24	3.75	3.69	**	+	**	+
30	HEMBA1004000	1.83	1.8	1.37	4.32	4.14	3.96	2.63	3.55	2.34	**	+	*	+
	HEMBA1004006	1.37	0.24	1.22	2.13	0.94	1.32	1.17	2.2	0.95				
	HEMBA1004007	6.04	2.39	4.27	10.96	12.86	11.45	4.4	6.87	6.82	**	+		
	HEMBA1004010	2.94	1.19	1.2	2.7	3.56	3.85	6.4	6.08	6.71		**	+	
	HEMBA1004011	1.7	0.78	0.96	2.15	2.36	1.93	1.31	2.48	1.53	*	+		
35	HEMBA1004012	3.28	1.3	2.31	6.01	4.99	6.51	2.69	3.48	2.84	**	+		
	HEMBA1004015	2.75	2.05	2.56	5.11	5.22	4.78	3.43	3.89	3.43	**	+	*	+
	HEMBA1004024	5.55	4.27	3.76	12.33	16.73	14.13	7.3	6.98	6.78	**	+	*	+
	HEMBA1004029	4.41	3.27	3.73	8.08	10.91	6.74	3.3	6.61	3.72	*	+		
	HEMBA1004038	2.95	1.3	1.87	2.87	2.27	1.5	1.15	4.52	1.31				
	HEMBA1004042	0.98	0.07	0.48	1.39	1.07	0.6	0.74	2.16	0.79				
40	HEMBA1004045	1.3	0.85	0.68	0.88	3.2	1.57	1.01	1.7	0.88				
	HEMBA1004048	7.55	3.12	4.4	8.61	11.26	7.66	7.55	7.12	7.08				
	HEMBA1004049	1.17	0.64	0.7	2.26	3.05	2.36	1.86	2.62	1.77	**	+	*	+
	HEMBA1004051	4.38	1.98	1.73	3.51	4.36	4.02	9.79	8.74	8.15		**	+	
	HEMBA1004053	8.83	5.44	4.46	17.89	9.59	13.59	14.13	13.15	12.39		**	+	
45	HEMBA1004055	2.65	0.36	1.81	2.57	2.89	2.7	1.3	3	1.6				
	HEMBA1004056	7.5	3.93	5.65	20.02	18.97	17.23	8.27	9.56	8.48	**	+	*	+
	HEMBA1004060	0.07	0.43	1.07	1.82	1.29	1.56	0.47	1.74	1.31	*	+		
	HEMBA1004061	14.25	5.05	4.22	4.44	4.07	3.17	2.96	3.91	3.1				
	HEMBA1004067	9.19	5.05	5.06	7.71	7.13	7.12	4.94	6.2	7.71				
50	HEMBA1004071	14.49	9.52	7.51	14.12	12.05	13.02	8.33	9.62	10.66				
	HEMBA1004074	7.06	2.77	2.38	5.08	5.28	4.16	4.21	4.55	5.63				
	HEMBA1004078	11.34	10.72	8	8.83	11.03	9.34	8.26	9.99	8.73				
	HEMBA1004085	3.75	2.6	2.94	3.51	4.46	4.01	3.83	4.11	2.49				
	HEMBA1004086	9.29	6.04	5.26	10.09	10	10.42	3.43	2.6	4.21				
55	HEMBA1004097	2.9	2.64	1.85	1.78	5.31	3.52	1.65	4.02	2.63				
	HEMBA1004100	5.05	2.67	3.16	5.99	4.86	5.28	5.44	5.79	5.01				
	HEMBA1004103	10.13	4.33	3.51	10.84	11.41	10.36	6.76	6.57	8.64				

Table 187

	HEMBA1003666	1.38	1.05	0.83	1.72	1.7	1.06	0.87	1.13	0.89				
	HEMBA1003667	14.71	11.01	9.94	14.75	22.82	18.78	15.24	10.59	14.49				
5	HEMBA1003670	0.91	0.22	0.29	1.11	1.61	1.82	0.56	1.43	0.85	*	+		
	HEMBA1003674	26.03	18.94	18.61	21.67	28.7	30.08	14.76	19.25	20.49				
	HEMBA1003677	3.73	1.52	2.36	7.63	8.16	6.96	10.74	10.88	8.28	**	+	**	+
	HEMBA1003679	1.48	0.67	1.25	5.41	5.58	4.44	1.61	3.27	2.24	**	+		
	HEMBA1003680	6.18	3.86	3.32	4.89	3.65	4.22	2.45	3.41	4.34				
10	HEMBA1003684	3.07	3.42	2.52	4.93	3.87	3.53	2.61	2.37	4.26				
	HEMBA1003690	8.67	4.5	4.89	6.53	5.61	6.33	6.11	7.01	7.57				
	HEMBA1003692	6.51	4.39	2.76	7.65	13.21	11.37	6.71	7.24	6.09	*	+		
	HEMBA1003702	7.49	3.3	2.54	5.23	6.69	4.84	4.77	3.72	5.73				
	HEMBA1003711	5.86	2.58	3.21	3.28	5.33	5.99	2.95	4.08	4.68				
15	HEMBA1003714	4.3	2.42	1.47	3.54	3.98	3.51	1.5	2.8	3.08				
	HEMBA1003715	5.16	2.24	2.94	8.09	8.13	8	2.66	4.48	4.1	**	+		
	HEMBA1003717	3.17	2.29	1.96	4.19	3.55	5.52	1.88	1.44	1.67	*	+		
	HEMBA1003720	1.56	1.73	1.27	3.11	3.53	3.49	2.3	1.66	3.08	**	+		
	HEMBA1003725	1.46	0.94	0.92	3.84	2.37	2.61	2.1	1.7	2.25	*	+	*	+
20	HEMBA1003728	6.2	3.24	4.06	5.16	6.27	6.67	5.85	4.48	3.55				
	HEMBA1003729	3.99	1.42	2.32	6.36	5.84	4.38	3.64	4.72	3.3	*	+		
	HEMBA1003732	1.63	1.1	1	3.52	2.12	1.25	0.95	1.54	1.47				
	HEMBA1003733	2.5	4.71	1.16	4.86	6.33	5.47	2.99	3.73	3.5				
	HEMBA1003742	6.12	2.9	4.2	5.24	4.87	5.32	2.62	6.27	5.03				
25	HEMBA1003743	2.64	1.63	1.2	2.32	2.37	3.69	2.34	1.46	1.92				
	HEMBA1003758	5.8	2.98	4.74	10.06	11.45	11.44	7.34	3.52	6.11	**	+		
	HEMBA1003760	5.32	2.29	2.62	4.55	3.7	4.58	3.57	4.5	4.37				
	HEMBA1003764	5.57	1.67	3.47	5.12	2.71	2.62	3.98	3.91	4.87				
	HEMBA1003769	11.09	7.81	6.22	7.38	7.99	10.09	8.32	6.25	8.19				
30	HEMBA1003773	4.06	2.15	2.74	3.4	2.78	2.89	3.34	3.66	3.74				
	HEMBA1003783	5.9	5.63	5.15	7.21	10.97	7.92	4.02	5.97	4.35				
	HEMBA1003784	1.56	0.55	0.26	1.01	1.64	1.14	0.84	1.59	1				
	HEMBA1003794	22.02	14.74	15.29	16.32	23.57	18.51	19.15	20.16	23.83				
	HEMBA1003799	3.18	0.83	0.69	1.6	1.44	2.62	1.76	1.29	1.44				
35	HEMBA1003803	5.18	3.99	2.9	7.41	7.07	8.96	7.06	6.07	5.89	*	+	*	+
	HEMBA1003804	4.31	3.24	3.27	5.11	3.19	3.64	3.81	3.96	3.99				
	HEMBA1003805	9.07	8.11	9.22	15.23	14.63	10.98	7.34	10.04	6.52	*	+		
	HEMBA1003807	2.26	0.57	1.05	1.41	1.99	1.42	1.14	1.69	1.19				
	HEMBA1003810	2.67	2.32	0.99	3.03	2.59	2.69	2.81	4.61	3.57				
40	HEMBA1003827	25.92	18.96	19.46	14.46	20.55	25.66	14.02	29.91	19.07				
	HEMBA1003836	9.8	5.94	7.41	16.46	20.73	18.84	10.1	7.05	9.1	**	+		
	HEMBA1003838	29.21	22.41	20.25	35.45	47.13	35.6	26.78	26.74	23.31	*	+		
	HEMBA1003843	8.31	5.73	4.45	4.63	2.15	3.75	3.72	2.7	3.17				
	HEMBA1003846	26.28	20.72	18.37	21.86	22.27	12.11	9.99	15.1	13.9		*	-	
	HEMBA1003856	3.23	2.48	1.56	1.62	2.7	2.03	1.6	2.63	2.09				
45	HEMBA1003857	5.6	3.94	4.15	8.14	11.16	11.16	4.4	7.01	4.61	**	+		
	HEMBA1003864	4.85	1.81	2.77	3.23	4.07	4.12	3.31	3.39	2.74				
	HEMBA1003866	1.42	0.62	1.37	1.22	1.21	1.69	0.7	1.76	0.89				
	HEMBA1003868	13.28	7.75	6.42	9.42	7.15	9.18	5.91	7.86	7.2				
	HEMBA1003879	2.14	1.42	1.52	4.08	4.35	3.36	3.08	2.74	2.28	**	+	*	+
50	HEMBA1003880	4.68	2.16	2.83	4.05	4.87	3.64	3.32	3.39	3.6				
	HEMBA1003884	5.74	3.71	3.92	4	4.49	5.17	4.44	4.54	4.54				
	HEMBA1003885	10.32	6.22	7.2	14.27	16.75	16.47	8.29	8.14	8.72	**	+		
	HEMBA1003887	5.7	2.76	3.69	4.8	4.75	6.26	4.28	5.31	3.98				
	HEMBA1003890	5.76	2.14	5.85	3.21	4.35	4.1	2.43	1.67	2.34				
55	HEMBA1003893	24.48	11.91	15.58	30.06	36.65	30.84	15.38	16.42	17.2	*	+		
	HEMBA1003896	19.51	13.33	11.04	17.41	24	18.91	15.03	15.04	20.6				
	HEMBA1003902	8.4	6.56	5.19	9.58	9.98	7.89	5.71	4.62	5.77				

Table 186

	HEMBA1003433	2.51	1.64	1.17	2.63	2.77	1.5	2.03	2.04	0.74				
5	HEMBA1003440	7.38	4.95	3.98	3.59	4.49	2.94	11.67	10.24	9.89		*	+	
	HEMBA1003442	7.11	3.89	5.36	33.69	44.16	39.43	12.88	14.11	14.92	**	+	**	+
	HEMBA1003447	6.43	2.84	5.38	2.86	4.59	3.43	2.19	3.65	2.78				
	HEMBA1003453	5.3	2.06	4.2	3.35	2.95	3.68	3.79	4.22	4.22				
	HEMBA1003461	4.9	1.85	2.53	3.24	4.51	4.52	2.91	4.48	2.29				
10	HEMBA1003463	2.07	0.69	1.15	5.59	5.7	5.89	4.6	5.83	5.74	**	+	**	+
	HEMBA1003465	9.37	4.59	4.46	10.69	9.03	7.99	6.08	6.86	6.92				
	HEMBA1003480	9.33	5.04	6.92	12.74	16.03	14.45	6.27	6.32	7.43	**	+		
	HEMBA1003485	20.75	10.29	10.54	10.17	12.27	12.15	10.87	6.69	7.13				
	HEMBA1003487	4.58	2.05	1.61	2.41	3.47	2.58	3.04	3.53	2.9				
	HEMBA1003492	2.07	1.37	0.95	2.53	2.7	2.94	1.03	2.89	1.4	*	+		
15	HEMBA1003494	2.49	0.76	1.49	27.92	31.78	20.12	3.6	6.11	5.48	**	+	*	+
	HEMBA1003497	3.12	0.78	1.83	3.69	4.28	3.96	1.74	2.6	2.31	*	+		
	HEMBA1003503	3.45	2.06	1.43	3.15	2.26	2.25	1.52	3	3.05				
	HEMBA1003511	2.69	1.04	0.98	1.76	1.46	1.83	1.71	1.33	0.95				
20	HEMBA1003528	18.14	11.27	11.45	12.37	19.83	18.44	16.97	12.4	16.79				
	HEMBA1003530	2.6	1.44	2.11	2.26	2.64	3.14	2.32	2.96	3.27				
	HEMBA1003531	6.99	4.57	4.74	10.98	15.62	10.36	6.08	6.8	4.37	*	+		
	HEMBA1003532	13.93	5.28	9.84	12.79	13.95	12.42	7.71	9.02	10.58				
	HEMBA1003538	2.36	1.42	1.55	0.71	3.61	2.87	1.32	3.05	1.48				
	HEMBA1003545	1.41	0.47	0.87	1.63	1.67	1.35	0.85	1.8	0.86				
25	HEMBA1003546	6.22	3.88	2.1	11.53	13.41	10.1	6.93	7.89	5.98	**	+		
	HEMBA1003548	0.92	0.44	0.29	1.8	1.25	1.92	0.41	1.43	0.38	*	+		
	HEMBA1003553	10.98	8.66	9.18	19.1	13.8	21.91	7.81	8.18	9.02	*	+		
	HEMBA1003555	3.02	1.7	1.46	1.76	3.2	2.69	2.27	3.4	2.27				
	HEMBA1003556	4.32	1.68	2.2	3.83	6.46	5.67	2.71	3.54	2.22				
30	HEMBA1003560	1.14	1.46	1.03	0.88	1.35	1.08	1.46	2.03	1.63				
	HEMBA1003565	4.06	3.07	3.95	3.82	4.6	4.62	4.01	5.7	5.12				
	HEMBA1003568	2.91	0.76	1.15	1.22	1.08	1.38	1.05	1.98	0.77				
	HEMBA1003569	8.99	12.88	9.75	5.29	6.55	5.16	4.54	5.33	5.68	*	-	*	-
	HEMBA1003571	10.48	4.42	3.13	21.11	11.99	10.73	5.96	8.94	7.11				
35	HEMBA1003579	5.23	2.72	1.87	4.14	3.57	5.4	3.01	3.4	2.79				
	HEMBA1003580	11.03	7.36	6.64	6.54	6.56	8.11	7.97	8.17	8.81				
	HEMBA1003581	5.6	4.24	4.26	4.68	5.52	5.87	5.47	4.38	5.02				
	HEMBA1003591	39.81	31.07	28.74	52.34	52.04	48.99	14.34	10.05	14.79	**	+	**	-
	HEMBA1003595	1.99	0.8	1.07	3.33	4.04	3.39	2.08	3.22	1.45	**	+		
40	HEMBA1003597	1.33	0.63	1.33	3.65	3.35	4.52	1.94	2.9	2.36	**	+	*	+
	HEMBA1003598	2.9	0.82	1.41	1.32	2.05	2.83	1.88	0.98	0.49				
	HEMBA1003600	5.78	3.55	3.06	6.44	7.48	5.87	4.2	4.07	5.87				
	HEMBA1003602	2.69	1.98	1.66	3.29	2.76	2.29	1.48	2.11	2.34				
	HEMBA1003604	11.43	8.02	8.72	12.24	9.01	11.87	7.65	8.32	8.25				
45	HEMBA1003610	8.44	6.02	5.83	14.76	14.29	15.88	13.42	9.31	12.15	**	+	*	+
	HEMBA1003615	6.42	3.45	3.87	5.96	5.91	5.22	3.28	5.75	3.48				
	HEMBA1003617	3.99	3.24	3.91	16.74	14.07	12.64	7.57	9.08	9.03	**	+	**	+
	HEMBA1003620	5.35	2.63	3.62	8.39	6.31	6.44	4.6	5.32	5.6	*	+		
	HEMBA1003621	5.01	4.74	3.02	9.46	12.07	10.28	5.9	5.67	5.82	**	+		
	HEMBA1003622	1.74	1.02	0.61	2.09	2.03	2.5	0.94	1.66	0.88	*	+		
50	HEMBA1003630	1.59	0.33	0.75	1.41	1.11	1.15	1.54	2.32	1.54				
	HEMBA1003637	2.15	0.95	0.99	3.26	5.54	3.57	1.75	2.71	1.99	*	+		
	HEMBA1003640	2.27	1.59	2.11	4.99	4.15	5.22	3.36	5.53	2.27	**	+		
	HEMBA1003645	1.63	0.53	1.13	3.97	2.86	2.71	2.66	3.44	1.36	*	+		
	HEMBA1003646	0.89	0.8	1.19	3.33	3.36	4.74	1.35	3.89	1.8	**	+		
55	HEMBA1003647	0.79	0.36	0.72	3.69	2.19	3.35	1.03	2.87	1.04	**	+		
	HEMBA1003656	3.32	1.76	1.62	4	4.27	4.72	3.61	3.92	2.65	*	+		
	HEMBA1003662	2.77	1.1	0.73	3.91	3.34	1.69	3.38	3	3.35				

Table 185

	HEMBA1003204	4.47	2.88	1.95	6.42	9.31	7.19	4.16	4.11	3.52	*	+		
	HEMBA1003210	5.3	3.32	2.57	8.14	10.48	8.37	17.93	9.12	16.05	**	+	*	+
5	HEMBA1003212	10.06	5.15	6.84	12.91	14.8	18.49	9.01	8.88	9.76	*	+		
	HEMBA1003218	1.85	0.63	1.04	1.36	1.25	1.72	1.4	2.5	1.89				
	HEMBA1003220	27.66	24	25.44	26.62	36.09	37.79	16.07	14.85	17.5		**	-	
	HEMBA1003222	2.88	1.72	3.36	3.75	3.58	3.59	2.57	3.59	2.87				
	HEMBA1003225	2.92	1.48	2.59	3.07	2.81	2.57	2.42	3.81	3.48				
10	HEMBA1003229	3.63	1	0.92	4.49	4.02	4.36	4.86	6.18	8.35		*	+	
	HEMBA1003230	4.81	1.33	1.59	3.63	3.48	2.96	4.65	4.11	4.45				
	HEMBA1003235	4.25	2.83	2.72	4.77	5.98	6.44	3.15	3.65	2.94	*	+		
	HEMBA1003236	2.61	2.12	2.62	4.85	3.24	5.32	5.66	4.6	3.8	*	+	*	+
	HEMBA1003250	1.73	0.34	1.4	2.93	3	2.03	1.83	2.23	1.38	*	+		
15	HEMBA1003252	5.88	2.96	5.36	7.78	7.79	7.89	4.58	5.63	5.99	*	+		
	HEMBA1003257	4.93	1.49	3.88	3.03	4.82	4.08	2.99	3.59	4.04				
	HEMBA1003268	0.75	0.26	0.6	2.39	1.18	1.2	0.42	1.31	0.41				
	HEMBA1003273	3.46	2.51	1.67	5.94	6.01	5.04	2.19	3.45	3.47	**	+		
	HEMBA1003276	1.81	1.29	0.96	4.38	4.69	4.83	2.14	2.73	3.03	**	+	*	+
20	HEMBA1003277	2.81	1.68	0.99	2.39	2.91	2.66	2.69	2.74	1.67				
	HEMBA1003278	1.65	0.9	1.98	2.98	3.92	3.95	2.37	3.01	2.17	**	+		
	HEMBA1003280	3.32	1.78	3	4.76	3.3	3.57	2.93	5.18	3.65				
	HEMBA1003281	4.06	0.91	2.42	3.46	3.32	3.57	2.53	4.81	3.88				
	HEMBA1003284	0.48	0.51	0.58	2.22	0.82	1.41	1.13	2.8	1.15				
25	HEMBA1003286	3.88	2.4	2.73	5.92	3.88	3.67	2.08	4.79	3.77				
	HEMBA1003291	2.38	1.74	0.96	2.57	3.95	3.8	2.72	4.5	5.97	*	+		
	HEMBA1003294	5.2	3.14	3.02	8.15	7.24	7.54	4.43	4.64	6.12	**	+		
	HEMBA1003296	3.52	1.49	1.47	1.62	2.44	1.83	2.01	2.49	2.5				
	HEMBA1003304	1.33	0.87	0.46	1.14	1.8	1.15	0.92	1.05	1.59				
30	HEMBA1003306	4.82	1.91	2.68	6.16	5.24	6.21	5.8	5.67	6.01	*	+	*	+
	HEMBA1003309	0.64	0.18	0.98	3.28	3.28	2.43	1.17	2.04	1.94	**	+	*	+
	HEMBA1003314	30.47	18.15	16.33	19.29	25.08	19.75	20.31	20.79	24.11				
	HEMBA1003315	10.03	5	5.86	8.82	6.71	8.85	7.02	6.3	8.18				
	HEMBA1003322	6.46	2.81	4.38	11.92	11.23	7.71	5.2	4.77	6.83	*	+		
	HEMBA1003326	4.18	1.78	2.35	2.75	2.35	2.84	2.28	3.1	3.12				
35	HEMBA1003327	1.82	3.14	1.29	2.95	3.45	3.27	2.29	2.03	3.08				
	HEMBA1003328	4.01	4	2.1	5.29	8.03	6.1	3.75	5.53	3.53	*	+		
	HEMBA1003330	11.21	6.43	6.46	11.55	10.31	11.11	5.39	5.56	6.86				
	HEMBA1003348	5.75	4.37	3.56	10.47	9.44	9.51	4.42	5.42	4.82	**	+		
	HEMBA1003349	3.52	2.39	2.06	5.95	6.68	6.94	3.15	4.91	3.36	**	+		
40	HEMBA1003370	20.51	11.56	11.02	25.15	23.1	21.13	12.45	14.72	17.99				
	HEMBA1003373	3.04	1.4	0.86	3.17	2.01	3.32	2.12	1.4	2.16				
	HEMBA1003376	11.18	5.54	7.92	20.96	23.88	21.25	10.64	10.28	11.12	**	+		
	HEMBA1003380	2.3	1.46	1.33	2.34	1.63	1.87	2.49	1.54	2.46				
	HEMBA1003384	2.29	0.73	1.56	3.93	3.22	3.27	1.72	2.42	3.12	*	+		
45	HEMBA1003387	1.34	0.55	1.92	1.88	0.47	0.99	1.2	0.99	1.1				
	HEMBA1003392	8.27	4.38	5.55	5.24	7.99	8.63	5.42	7.97	6.53				
	HEMBA1003395	1.96	1.22	1.19	2.43	3.54	3.02	1.59	5.5	1.02	*	+		
	HEMBA1003399	5.58	3.74	3.33	5.08	4.37	5.04	3.4	3.1	3.67				
	HEMBA1003400	10.74	5.28	6.5	8.13	8.07	5.69	7.43	7.79	7.28				
50	HEMBA1003402	4.66	2.07	1.57	4.25	3.02	2.77	2.27	1.71	2.18				
	HEMBA1003403	4.57	4.91	4.99	4.14	4.8	3.26	2.96	3.55	2.8		**	-	
	HEMBA1003408	10.68	7.13	5.44	7.16	7.17	7.67	7.62	9.08	7.52				
	HEMBA1003412	6.57	4.94	4.07	6.42	6.27	6.69	3.99	6.63	4.67				
	HEMBA1003417	4.27	2.26	3.09	1.9	2.03	2.19	2.24	2.99	2.76				
55	HEMBA1003418	10.03	4.9	6.22	10.24	12.15	12.3	3.46	6.64	4.53				
	HEMBA1003420	1.52	0.53	0.73	7.04	2.2	5.31	11.33	12.68	10.88		**	+	
	HEMBA1003425	1.37	1.11	1.09	2.68	2.01	2.17	1.69	0.88	0.88	**	+		

Table 184

	HEMBA1002992	9.81	4.26	5.48	8.62	8.82	10.46	6.24	6.81	7.68				
	HEMBA1002995	9.95	5.67	5.79	12.67	13.82	15.45	7.42	8.68	5.34	*	+		
5	HEMBA1002997	5.35	3.23	2.63	6.04	6.82	4.47	3.67	4.27	4.14				
	HEMBA1002999	1.41	1.2	1	1.77	1.86	2.15	1.32	1.54	1.79	*	+		
	HEMBA1003004	4.4	2.05	2.04	4.44	2.35	3.6	4.34	2.86	3.43				
	HEMBA1003006	3.81	3.03	1.95	4.39	5.85	4.42	3.51	4.58	4.16				
	HEMBA1003008	3.21	2.19	2.5	3.68	6.17	6.62	2.11	3.8	2.8	*	+		
10	HEMBA1003021	7.74	5.2	3.87	9.69	18.49	13.68	7.18	7.68	6.46	*	+		
	HEMBA1003027	2.46	2.25	2.2	3.48	3.21	5.26	3.71	4.99	5.27		**	+	
	HEMBA1003029	16.49	15.58	12.66	14.01	22.6	13.51	9.84	22.76	21.22				
	HEMBA1003031	7	6.8	4.83	11.72	14.51	12.51	5.21	5.97	6.1	**	+		
	HEMBA1003032	8.54	5.52	5.51	6.83	9.05	7.67	7.01	6.8	7.13				
15	HEMBA1003033	13.69	8.92	7.92	18.19	20.22	19.59	7.06	10.97	9.51	**	+		
	HEMBA1003034	10.16	6.76	5.59	16.34	16.21	18.88	7.61	9.38	7.94	**	+		
	HEMBA1003035	0.86	0.59	0.52	1.61	1.97	0.55	0.09	2.49	0.47				
	HEMBA1003037	14.14	5.43	5.96	7.58	8.71	8.97	7.73	6.56	7.19				
	HEMBA1003041	13.54	5.42	7.39	11.23	11.7	11.68	7.62	7.38	7.89				
20	HEMBA1003046	10.88	8	7.65	10.34	12.65	9.57	6.83	7.13	6.72				
	HEMBA1003047	6.06	2.52	2.2	4.15	5.03	5.74	4.14	4.89	5.07				
	HEMBA1003048	4.06	2.13	2.64	5.2	6.24	5.07	5.54	7.31	7.12	*	+	**	+
	HEMBA1003064	1.85	0.88	1.11	2.44	3.01	2.83	0.75	2.55	1.61	*	+		
	HEMBA1003067	3.99	3.75	3.24	6.04	4.55	5.67	2.42	3.41	2.73	*	+		
25	HEMBA1003071	4.89	2	2.46	3.09	3.36	3.95	2.75	4.15	2.46				
	HEMBA1003072	5	3.54	3.49	9.31	7.84	7.21	4.62	3.3	3.28	**	+		
	HEMBA1003076	17.78	7.65	8.23	14.31	15.74	15.39	9.87	12.41	9.92				
	HEMBA1003077	2.58	1.45	1.89	1.93	2.25	1.91	1.66	2.15	1.57				
	HEMBA1003078	2.54	1.51	1.55	3	4.23	3.44	2.66	2.38	2.18	*	+		
	HEMBA1003079	1.91	1.85	1.65	2.48	2.95	4.35	2.42	3.49	4.71				
30	HEMBA1003083	3.9	3.64	3.64	4.53	10.29	6.35	3.33	5.25	3.51				
	HEMBA1003086	4.22	1.35	2.59	5.79	6.56	7.76	2.96	3.81	3.22	*	+		
	HEMBA1003090	4.24	1.39	3.28	3.62	3.38	4	2.6	2.81	2.75				
	HEMBA1003094	7.91	4.48	3.84	6.39	5.65	6.37	3.63	7.81	6.29				
35	HEMBA1003096	2.55	1.26	1.42	4.86	4.02	4.86	6.22	5.89	7.19	**	+	**	+
	HEMBA1003098	13.3	7.22	6.89	14.21	8.08	14.42	7.57	4.87	3.41				
	HEMBA1003101	3.86	1.83	2.21	3.42	3.01	2.84	5.05	3.05	3.57				
	HEMBA1003109	4.5	2.81	2.78	4.25	4.4	4.14	3.32	3.04	3.75				
	HEMBA1003114	4.72	1.49	2.76	4.85	5.37	3.83	2.72	3.02	2.47				
	HEMBA1003117	3.34	1.32	1.84	1.94	3.48	3.4	1.15	2.8	1.47				
40	HEMBA1003120	6.26	3.04	4.46	8.53	11.03	9.87	2.19	4.52	4.23	*	+		
	HEMBA1003129	2.92	2.45	1.66	4.57	4.93	4.76	2.42	2.95	3.22	**	+		
	HEMBA1003133	3.76	2.75	2.66	4.15	4.94	3.64	3.03	3.83	3.17				
	HEMBA1003136	10.1	5.38	5.56	4.76	5.69	3.55	4.84	6.06	5.16				
	HEMBA1003142	3.63	2.31	2.57	5.7	6.12	5.75	4.06	4.52	4.11	**	+	*	+
45	HEMBA1003148	3.76	1.84	1.85	6.57	6.35	7.13	3.78	3.2	3.41	**	+		
	HEMBA1003151	3.06	1.21	2.06	3.57	3.12	3.47	1.14	2.71	1.88				
	HEMBA1003152	0.94	1.17	1.24	1.37	1.78	3.11	1.18	1.39	0.96				
	HEMBA1003157	5.21	1.69	3.38	3.87	2.5	3.42	0.86	1.61	0.57				
	HEMBA1003166	16.26	12.43	11.19	32.39	35.71	31.69	16.79	24.31	19.76	**	+		
50	HEMBA1003171	2.89	0.72	0.57	1.31	1.88	0.92	1.38	1.84	1.16				
	HEMBA1003175	2.6	1.51	1.44	3.64	3.88	4.37	1.98	2.4	2.06	**	+		
	HEMBA1003179	4.43	2.72	4.24	3.15	2.78	4.29	3.01	4.28	3.14				
	HEMBA1003186	8.23	6.45	5.52	11.84	15.62	13.23	6.12	9.01	7.56	**	+		
	HEMBA1003196	5.41	2.8	3.15	4.6	6.06	5.01	3.65	4.6	3.58				
55	HEMBA1003197	1.16	0.72	0.59	1.88	2.27	1.84	1.37	1.38	0.69	**	+		
	HEMBA1003199	2.2	0.82	0.97	3.9	3.83	3.59	1.11	2.24	1.33	**	+		
	HEMBA1003202	6.51	4.3	4.72	9.3	10.18	10.03	5.76	4.86	4.67	**	+		

Table 183

	HEMBA1002716	2.33	1.79	1.1	2.97	1.95	2.33	1.67	1.27	1.06			
	HEMBA1002718	16.72	11.81	9.31	17.97	12.98	15.17	10.44	10.19	11.13			
5	HEMBA1002728	9.67	3.54	5.97	10.6	12.96	15.33	7.76	5.08	9.8	*	+	
	HEMBA1002730	7.86	2.52	3.4	5.36	7.91	6.74	6.78	4.96	7.37			
	HEMBA1002734	7.73	4.31	3.55	7.93	6.46	7.46	5.62	6.29	6.83			
	HEMBA1002742	3.65	1.6	2.01	2.64	2.74	2.6	1.48	2.29	1.92			
	HEMBA1002746	6.82	4.06	4.19	4.98	4.66	5.4	2.78	4.2	3.33			
10	HEMBA1002748	4.16	2.16	3.32	2.53	4.45	4.03	2.42	2.8	3.88			
	HEMBA1002750	6.45	3.44	3.09	5.38	6.22	7.28	3.44	2.24	3.97			
	HEMBA1002755	6.83	3.3	3.88	9.75	9.18	10.07	4.45	5.29	5.42	*	+	
	HEMBA1002759	2.47	0.92	1.55	4.32	3.79	4.12	2.56	2.66	2.65	**	+	
	HEMBA1002763	17.79	8.69	9.49	11.93	10.68	12.89	9.46	10.98	11.31			
15	HEMBA1002767	4.86	3.64	4.15	4.69	4.37	5.27	4.84	4.88	6.63			
	HEMBA1002768	7.65	3.89	4.38	7.6	7.36	7.85	6.31	6.75	7.16			
	HEMBA1002769	6.55	2.6	4.29	4.3	5.76	5.49	4.08	5.08	4.57			
	HEMBA1002770	10.29	6.74	8.19	11.22	11.06	13.4	7.03	6.36	8.42			
	HEMBA1002777	9.75	4.7	5.71	8.59	8.79	9.46	6.38	4.3	7.48			
20	HEMBA1002779	19.22	10.66	6.22	15.16	13.63	10.21	10.37	10.01	10.31			
	HEMBA1002780	5.7	2.86	3	6.99	7.8	9.55	4.79	4.73	6.4	*	+	
	HEMBA1002790	4.99	2.33	3.07	6.37	8.93	7.96	4.08	3.78	4.9	*	+	
	HEMBA1002794	8.37	5.67	4.58	5.78	6.13	8.44	6.79	6.5	6.15			
	HEMBA1002798	1.26	0.86	1.65	2.72	2.3	1.86	0.87	2.64	0.77	*	+	
25	HEMBA1002801	1.99	0.93	1.36	4.21	3.6	1.85	2.71	2.29	3.22		*	+
	HEMBA1002810	9.65	4.37	5.68	13.26	12.11	9.75	5.27	6.41	6.28			
	HEMBA1002816	9.84	4.52	4.72	9.31	6.58	9.2	5.89	5.54	5.86			
	HEMBA1002818	13.95	7.65	7.85	12.57	11.48	11.5	11.94	8.46	10.87			
	HEMBA1002820	8.63	4.01	5.8	12.08	16.06	13.75	7.38	6.93	7.73	*	+	
30	HEMBA1002826	2.06	0.77	0.96	1	0.94	1.69	1.3	2.13	0.88			
	HEMBA1002833	9.88	4.57	5.73	7.08	7.89	7.35	7.95	8.57	7.16			
	HEMBA1002850	0.76	0.3	1.24	1.8	1.57	1.81	0.67	2.12	1.24	*	+	
	HEMBA1002862	2.92	2.24	3.55	9.63	8.86	7.72	5.29	8.86	7.89	**	+	+
	HEMBA1002863	3.16	2.79	5.23	4.86	5.55	5.31	3.6	5.86	5.95			
35	HEMBA1002867	3.74	1.09	1.41	1.95	2.42	2.24	1.51	1.85	1.96			
	HEMBA1002876	10.81	3.46	4.85	5.22	5.51	6.47	5.11	4.45	5.37			
	HEMBA1002886	1.73	1.14	1.2	1.8	3.11	2.84	1.24	1.52	0.93	*	+	
	HEMBA1002896	5.56	2.89	2.26	4.16	5.6	6.36	4.43	4.26	5.28			
	HEMBA1002913	6.83	3.41	4.1	6.13	4.56	5.54	4.6	4.46	4.22			
	HEMBA1002921	5.09	1.35	3.42	4.01	3.76	3.47	2.82	3.68	1.76			
40	HEMBA1002924	3.44	1.46	2.03	3.99	2.79	5.07	4.7	2.86	2.66			
	HEMBA1002934	19.41	10.56	13.01	28.28	26.9	31.77	13.81	10.62	17.37	**	+	
	HEMBA1002935	5.64	2.51	3.1	9.39	9.17	8.78	4.05	4.5	6.44	**	+	
	HEMBA1002937	2.94	0.97	1.56	5.32	3.72	3.3	4.25	3.23	5.41		*	+
	HEMBA1002939	5.23	2.26	1.27	6.12	6.22	7.2	3.36	5.43	4.03	*	+	
45	HEMBA1002944	2.39	1.05	0.97	2.45	2.94	1.89	1.97	1.66	1.79			
	HEMBA1002951	4.82	2.48	2.82	6.08	7.02	6.04	4.67	6.63	5.8	*	+	
	HEMBA1002954	3.07	1.62	1.21	5.05	3.53	2.74	1.86	3.21	2.77			
	HEMBA1002962	4.7	4.71	2.06	11.63	8.54	7.28	2.97	4.52	4.25	*	+	
	HEMBA1002968	7.62	3.18	4.17	11.44	8.51	9.98	4.32	4.58	5.82	*	+	
50	HEMBA1002970	1.55	2.24	2.05	3.8	4.05	2.91	1.8	3.84	2.44	*	+	
	HEMBA1002971	2.55	2.17	1.09	2.11	2.8	2.2	1.84	1.44	2.55			
	HEMBA1002973	4.7	1.37	2.41	7.46	7.53	5.02	4.19	3.07	3.54	*	+	
	HEMBA1002978	4.6	2.2	2.96	5.07	6.26	5.9	2.87	3.98	1.74	*	+	
	HEMBA1002981	10.14	3.92	5.05	6.62	5.73	6.85	4.75	3.37	4.22			
55	HEMBA1002985	5.65	3.15	2.63	4.75	6.26	6.46	4.22	6.1	4.66			
	HEMBA1002986	8.06	6.02	4.91	10.22	16.35	15.33	12.26	16.22	12.17	*	+	+
	HEMBA1002988	1.58	0.97	1.43	5.23	7.34	7.21	3.78	3.98	3.2	**	+	**

Table 182

	HEMBA1002486	8.76	6.38	4.66	8.52	8.8	10.2	5.18	4.82	7				
5	HEMBA1002490	4.65	2.87	1.43	3.68	2.57	3.08	3.01	1.75	3.98				
	HEMBA1002495	3.72	2.75	1.63	4.11	3.81	4.48	2.24	3.36	3.9				
	HEMBA1002498	2.75	1.45	1.13	1.68	1.82	1.19	2.23	1.05	1.96				
	HEMBA1002501	4.03	2.44	2.73	2.79	3.44	4.73	2.7	2.56	4.15				
	HEMBA1002503	5.04	2.61	2.84	6.45	4.88	5.28	3.23	3.79	3.13				
10	HEMBA1002504	8.07	4.4	4.13	10.71	10.32	10.08	4.47	6.58	5.92	*	+		
	HEMBA1002508	5.99	4.98	4.38	8.82	14.4	16.34	4.8	6.77	5.33	*	+		
	HEMBA1002513	8.6	4.28	4.52	7.08	4.68	6.71	4.93	3.86	4.51				
	HEMBA1002515	4.33	1.73	2.07	3.29	2.16	3.66	2.65	1.63	3.58				
	HEMBA1002524	9.35	6	4.75	8.16	6.47	7.51	5.77	5.05	6.67				
	HEMBA1002538	4.58	2.05	1.84	2.98	3.05	4.53	2.16	2.92	2.68				
15	HEMBA1002542	8.07	5.4	5.41	9.41	8.04	9.27	4.65	5.75	5.16				
	HEMBA1002544	3.1	1.76	1.69	4.47	3.6	3.68	2.18	2.17	2.61	*	+		
	HEMBA1002546	50.52	34.29	29.94	56.51	60.33	61.14	35.34	44.64	38.68	*	+		
	HEMBA1002547	2.2	1.72	2.07	1.6	3.25	2.8	2.97	4.34	2.32				
	HEMBA1002550	7.14	5.4	3.96	4.54	4.38	4.87	6.51	4.38	5.24				
20	HEMBA1002551	5.47	2.09	2.27	5.04	4.39	3.41	4.06	3.2	3.87				
	HEMBA1002552	12.19	3.86	6.34	10.16	9.24	10.66	6.5	6.73	6.78				
	HEMBA1002555	1.98	0.86	1	1.95	2.49	2.76	2.25	1.97	2.82				
	HEMBA1002558	7.34	3.99	4.45	10.47	9.14	11.18	5.75	4.9	5.48	*	+		
	HEMBA1002561	1.53	2.23	1.45	3.76	4.16	3.85	2.34	2.9	2.42	**	+		
25	HEMBA1002562	2.58	1.09	1.24	1.55	1.58	1.46	1.13	1.38	1.77				
	HEMBA1002568	4.34	2.05	1.84	2.65	3.18	3.63	2.01	3.91	2.77				
	HEMBA1002569	10.12	2.96	3.15	6.04	6.91	7.8	6.66	5.49	5.73				
	HEMBA1002570	17.18	8.39	8.43	7.74	7.84	6.32	4.15	4.68	4.47				
	HEMBA1002574	9.13	5.2	4.08	4.71	4.69	3.46	6.41	4.34	4.04				
30	HEMBA1002583	2.63	1.94	1.44	4.35	4.76	4.81	4.07	4.23	4.71	**	+	**	+
	HEMBA1002587	9.65	5.73	4.29	5.38	5.09	6.69	6.95	4.55	5.87				
	HEMBA1002590	5	2.82	3.17	5.3	7.12	7.9	3.16	4.25	3.45	*	+		
	HEMBA1002592	7.22	3.8	5.73	9.2	7.27	11.07	4.7	6.52	5.38				
	HEMBA1002595	6.26	2.72	4.83	2.78	4.06	4.2	3.48	5.01	4.73				
35	HEMBA1002609	4.35	4.09	2.17	4.02	4.01	4.31	3.53	3.64	3.18				
	HEMBA1002617	3.95	2.7	1.65	11.81	11.46	11.36	4.49	2.86	3.96	**	+		
	HEMBA1002619	6.56	3.72	3.15	6.01	4.48	4.66	4.55	5.76	4.4				
	HEMBA1002621	1.33	2.05	0.58	1.87	2.25	1.68	1.25	2.13	1.22				
	HEMBA1002624	10.87	5.76	5.5	10.8	7.15	9.93	6.61	9.08	8.33				
40	HEMBA1002628	2.46	1.89	1.56	8.26	8.9	8.55	6.5	6.76	7.64	**	+	**	+
	HEMBA1002629	2.92	1.59	1.72	2.35	1.78	3.13	2.15	2.22	2.55				
	HEMBA1002632	3.01	3.25	2.45	3.55	4.56	5.56	2.28	3.32	2.22				
	HEMBA1002645	5.23	3.12	3.3	9.71	9.85	8.47	4.08	4.56	3.84	**	+		
	HEMBA1002651	2.74	3.3	3.7	3.63	3.14	3.96	3.35	3.43	3.18				
	HEMBA1002652	10.09	4.55	2.8	3.94	7.21	6.45	4.43	4.54	4.59				
45	HEMBA1002659	10	4.51	4.33	9.97	11.66	8.87	6.05	6.33	5.41				
	HEMBA1002661	4.42	2.54	1.79	6.87	7.34	8.27	4.97	3.82	3.85	**	+		
	HEMBA1002666	3.37	1.93	1.85	2.66	2.68	3.23	2.59	2.71	2.16				
	HEMBA1002667	3.38	2.19	2.12	5.95	5.18	5.02	0.91	3	2.2	**	+		
	HEMBA1002673	24.31	16.62	13.86	16.81	24.76	24.88	10.84	15.49	10.36				
50	HEMBA1002678	6.22	4.52	2.39	8.83	7	9.06	5.27	5.81	5.58	*	+		
	HEMBA1002679	6.14	2.98	2.3	7.06	5.91	5.92	4.9	4.03	4.55				
	HEMBA1002688	2.43	1.85	1.49	1.28	2.14	1.93	0.91	1.94	1.12				
	HEMBA1002696	5.94	3.2	2.68	4.46	5.16	4.1	3.48	3.38	3.28				
	HEMBA1002703	14.6	8	9.65	11.88	14.48	12.25	7.74	9.23	10.55				
55	HEMBA1002706	14.74	6.11	9.63	11.39	13.51	13.79	6.16	6.12	6.47				
	HEMBA1002712	5.57	3.22	3.91	7.46	9.02	7.13	2.62	3.73	3.7	*	+		
	HEMBA1002715	7.56	4.05	6.71	7.13	9.71	10.17	4.38	6.93	5.48				

Table 181

	HEMBA1002262	19.33	13.63	11.06	41.08	43.27	39.59	22.08	18	19.52	**	+		
	HEMBA1002265	5.77	2.24	2.87	4.81	4.22	4.54	5.24	3.12	3.12				
5	HEMBA1002267	6.66	4.16	4.1	9.3	9.47	10.3	6.16	5.48	5.21	**	+		
	HEMBA1002270	6.24	3.34	3.58	7.78	8.98	7.9	3.76	4.01	4.91	*	+		
	HEMBA1002286	2.71	2.63	1.38	2.66	4.03	3.95	2.71	3.54	3.82				
	HEMBA1002290	7.29	3.76	4.66	10.41	13.32	10.04	5.97	7.37	8.02	*	+		
	HEMBA1002302	11.09	4.74	4.9	14.47	14.16	14.79	5.34	6.14	6.47	*	+		
10	HEMBA1002304	2.15	1.99	3.2	4.13	2.57	4.4	1.76	2.42	1.31				
	HEMBA1002307	20.52	10.07	9.13	9.76	9.15	8.21	13.28	14.24	17.09				
	HEMBA1002316	21.96	17.53	15.62	14.66	14.2	13.67	14.54	18.03	17.67				
	HEMBA1002319	3.87	2.44	2.95	2.86	3.71	4.51	3	3.92	3.09				
	HEMBA1002320	2.67	1.82	1.12	4.11	5.01	6.24	3.84	4.14	3.6	*	+	*	+
15	HEMBA1002321	1.46	2.38	0.87	3.05	1.97	2.21	1.05	1.18	1.29				
	HEMBA1002328	4.66	1.71	1.99	5.92	5.51	4.89	3	3.99	2.69				
	HEMBA1002333	4.92	1.14	2.37	2.57	3.45	2.77	2.04	3.2	1.93				
	HEMBA1002337	5.38	3.22	4.87	9.22	12.3	11.34	4.19	5.44	4.11	**	+		
	HEMBA1002339	23.81	10.43	6.17	11.11	15.1	14.91	11.67	12.27	12.43				
20	HEMBA1002341	7.39	3.74	4.25	4.55	4.12	3.82	6.09	5.66	5.69				
	HEMBA1002348	2.07	1.83	0.9	1.44	1.88	2.08	1.92	2.6	1.34				
	HEMBA1002349	1.51	1.42	0.34	1.38	1.3	1.96	1.46	2.19	1.38				
	HEMBA1002353	1.79	1.25	2.28	2.64	3.11	3.43	2.11	1.34	1.36	*	+		
	HEMBA1002356	13.39	6.02	7.85	8.42	10.26	11	4.88	6.24	6.12				
25	HEMBA1002357	136.4	89.6	109	142.6	135.4	152.8	57.09	66.8	75.58		*	-	
	HEMBA1002360	6.54	3.66	5.93	10.16	10.44	10.51	8.07	9.62	8.15	**	+	*	+
	HEMBA1002363	9.05	6.26	4.11	8.4	5.32	7.47	3.78	3.67	4.84				
	HEMBA1002365	2.33	1.04	1.69	2.69	1.93	1.79	0.53	1.83	2.11				
	HEMBA1002370	2.04	0.84	0.68	5.63	6.49	6.21	1.4	3.02	2.46	**	+		
30	HEMBA1002374	8.05	4.75	3.85	6.96	7.96	4.55	6.91	5.19	7.37				
	HEMBA1002376	22.58	10.7	11.64	20.42	22.01	21.09	9.22	9.95	12.27				
	HEMBA1002377	22.23	20.26	24.74	17.13	16.56	16.97	12.65	5.84	13.5	*	-	*	-
	HEMBA1002380	10.33	4.73	6.12	25.3	20.75	23.1	10.39	11.3	10.43	**	+		
	HEMBA1002381	6.11	3.6	4.83	7.07	8.7	10.4	3.87	4.54	4.53	*	+		
35	HEMBA1002384	15.5	10.84	6.42	29.27	32.78	29.1	8.58	9.53	10.47	**	+		
	HEMBA1002389	4.27	1.82	1.04	3.34	2.49	2.48	1.75	2.27	2.21				
	HEMBA1002396	5.31	1.45	2.21	3.61	3.86	4.27	4.37	4.75	6.22				
	HEMBA1002402	4.83	1.75	1.81	2.54	2.69	3.67	3.46	2.38	3.41				
	HEMBA1002417	10.95	4.91	5.09	7.22	6.91	7.47	6.16	5.78	7.28				
40	HEMBA1002419	5.08	2.09	1.3	5.6	4.81	5.12	3.66	3.43	3.31				
	HEMBA1002420	9.17	4.99	7.48	15.98	15.18	16.1	7.55	7.5	9.4	**	+		
	HEMBA1002421	3.35	2.15	2.59	6.22	6.03	5.26	6.83	6.92	8.74	**	+	**	+
	HEMBA1002423	1.54	0.63	0.83	2.44	3.48	3.88	1.7	2.52	2.63	*	+	*	+
	HEMBA1002424	8.4	2.37	3.82	4.43	5.04	5.22	3.68	2.96	3.47				
45	HEMBA1002426	6.49	3.96	3.42	4.7	3.88	6.41	6.1	4.1	4.43				
	HEMBA1002430	2.26	0.52	0.37	1.48	0.95	2.01	1.47	2.19	1.34				
	HEMBA1002439	5.88	2.46	3.67	5.4	4.16	4.53	3.95	4.58	3.74				
	HEMBA1002441	9.17	5.14	6.07	34.35	24.41	25.8	23.81	16.3	21.56	**	+	**	+
	HEMBA1002454	5.79	2.67	3.42	5.87	3.7	3.81	3.97	3.14	3.78				
	HEMBA1002458	25.18	17.65	26.81	56.49	54.86	61.69	25.04	31.59	38.07	**	+		
50	HEMBA1002460	13.9	7.3	5.63	4.27	4.5	4.21	3.8	3.84	3.88				
	HEMBA1002462	5.97	3.49	2.63	4.68	5.48	5.1	5.76	6.26	5.62				
	HEMBA1002465	1.48	0.35	0.87	1.94	1.91	2	2	1.62	1.54	*	+		
	HEMBA1002469	10.61	5.54	6.1	9.43	9.29	9.35	6.49	6.37	7.65				
	HEMBA1002475	2.44	1.25	1.2	2.62	1.19	1.93	1.75	2.35	1.4				
55	HEMBA1002477	4.33	2.21	3.54	6.33	6.45	9.03	4.12	5.33	4.08	*	+		
	HEMBA1002480	12.76	7.21	8.4	13.9	9.22	9.97	6.6	9.31	9.9				
	HEMBA1002481	4.17	1.44	3.57	5.7	5.97	7.71	3.35	4.98	3.12	*	+		

Table 180

	HEMBA1002084	1.07	0.5	0.79	1.77	1.77	2.12	1.8	1.72	1.8	**	+	**	+
	HEMBA1002085	15.53	10.5	9.09	3.93	5.17	5.54	4.22	4.66	4.34	*	-	*	-
5	HEMBA1002092	6.36	2.95	3.86	3.82	3.84	2.97	3.77	3.66	5.02				
	HEMBA1002098	2.76	1.13	1.81	2.4	2.24	2.53	2.57	2.73	1.55				
	HEMBA1002100	32.5	21.44	18.67	25.5	28.16	25.35	18.35	13.17	17.71				
	HEMBA1002101	14.23	9.44	8.67	29.98	21.28	21.61	20.63	10.83	13.44	*	+		
	HEMBA1002102	5.78	2.45	5.61	10.26	9.25	10.76	5.53	7.91	7.68	**	+		
10	HEMBA1002105	3.54	2.37	3.22	6.12	5.06	5.65	3.82	6.51	5.09	**	+		
	HEMBA1002107	11.45	5.11	6.25	8.68	8.52	8.38	12.57	12.66	17.5				
	HEMBA1002113	32.25	19.17	17.4	39.34	45.35	45.81	28.29	21.95	34.31	*	+		
	HEMBA1002119	2.11	2.17	0.99	2.79	2.14	2.54	2.06	2.87	1.79				
	HEMBA1002125	5.95	2.4	2.92	5.45	9.25	7.16	7.44	6.34	6.72				
15	HEMBA1002131	5.93	2	3.14	4.14	4.06	4.13	3.5	4.3	3.28				
	HEMBA1002133	6.81	5.25	2.52	6.36	5.83	7.36	4.72	7.3	4.48				
	HEMBA1002139	1.09	0.26	0.36	1.2	0.84	1.33	0.99	2.43	0.56				
	HEMBA1002141	1.29	0.49	1.21	2.38	1.03	1.99	0.5	1.42	1.34				
	HEMBA1002144	5.69	3.1	2.06	7.29	6.78	8.63	2.59	3.43	5.33	*	+		
20	HEMBA1002147	21.38	10.63	10.33	16.26	8.66	14.72	7.7	9.8	14.04				
	HEMBA1002150	19.09	10.95	13.29	13.45	10.91	11.19	15.49	16.53	17.44				
	HEMBA1002151	5.57	4.52	3.73	5.15	5.43	4.75	6.45	4.35	4.86				
	HEMBA1002153	2.06	0.67	0.65	2.43	2.33	1.79	1.41	1.49	1.24				
	HEMBA1002156	6.64	2.07	2.79	3.49	2.76	4.92	4.24	4.29	3.26				
25	HEMBA1002160	9.96	4.66	4.52	11.03	12.78	11.54	5.12	4.86	6.62	*	+		
	HEMBA1002161	5.93	2.84	3.76	7.56	5.8	7.54	3.32	4.13	3.25				
	HEMBA1002162	7.92	3.54	4.29	9.23	12.27	9.59	6.96	4.68	6.43	*	+		
	HEMBA1002163	16.52	8.9	8.29	30.66	23.8	18.1	23.47	24.41	36.58	*	+	*	+
	HEMBA1002164	6.58	3.37	3.2	7.61	7.12	6.96	5.68	4.84	5.16				
30	HEMBA1002166	39.64	27.28	27.86	36.11	45.05	43.8	20.24	20.85	22.71				
	HEMBA1002167	4.76	1.86	1.62	2.99	2.78	2.27	3.13	3.05	2.27				
	HEMBA1002173	5.99	4.25	4.52	7.86	9.55	7.59	5.43	4.55	6.47	*	+		
	HEMBA1002177	7.43	2.78	2.92	3.23	3.61	5.94	3.11	3.88	4.09				
	HEMBA1002178	5.72	4.28	4.98	4.38	4.69	4.23	3.54	5.04	4.32				
	HEMBA1002179	38.56	31.74	22.53	17.89	19.71	18.71	27.72	23.97	26.16				
35	HEMBA1002185	6.54	3.16	3.12	9.32	10.15	8.6	6.14	5.78	6.76	*	+		
	HEMBA1002188	8.98	4.74	6.39	7.79	6.15	7.58	6.43	5.81	6.6				
	HEMBA1002189	3.48	3.26	1.78	4.27	5.47	4.09	2.69	3.88	3.54				
	HEMBA1002191	8.3	3.89	4.67	8.84	6.83	6.19	5.91	5.98	6.36				
	HEMBA1002192	5.28	4.26	4.29	8.27	6.01	5.9	2.94	2.49	2.82		**	-	
40	HEMBA1002195	5.98	3.67	4.11	6.21	5.77	4.89	3.93	4.26	3.98				
	HEMBA1002196	1.16	1.29	1.53	2.22	2.69	3.34	2.25	2.29	2.94	*	+	**	+
	HEMBA1002199	2.9	1.1	2.41	4.59	4.69	3.07	3.88	2.62	3.82				
	HEMBA1002204	3.61	1.66	0.98	2.22	2.66	1.99	3.47	1.11	1.87				
45	HEMBA1002208	48.26	35.92	30.61	48.99	56.44	45.32	18.77	22.83	23.91		*	-	
	HEMBA1002212	1.63	2.93	1.64	4.46	4.61	4.63	3.31	1.91	1.67	**	+		
	HEMBA1002215	6.24	3.92	3.6	5.45	4.91	5.62	4.3	4.83	3.3				
	HEMBA1002217	18.63	10.54	10.96	10.92	19.47	21.75	8.18	9.72	7.73				
	HEMBA1002220	2.36	1.42	1.13	2.73	2.21	2.69	1.63	2.43	2.05				
	HEMBA1002226	7.06	3.57	4.14	9.44	8.41	9.81	3.48	7.93	5.79	*	+		
50	HEMBA1002227	23.89	11.28	12.65	63.81	64.96	61.28	43.22	41.38	46.55	**	+	**	+
	HEMBA1002229	12.93	9.6	8.96	22.59	17.4	16.24	11.18	9.43	9.71	*	+		
	HEMBA1002237	2.73	1.56	1.22	2.88	3.54	2.57	2.65	2.19	1.5				
	HEMBA1002239	9.11	4.97	3.45	14.42	9.61	11	4.46	5.84	6.91				
	HEMBA1002241	4.16	2.92	3.35	4.29	3.16	4.98	3.45	3.5	3.33				
55	HEMBA1002253	2	1.21	0.86	1.18	1.75	2.13	1.77	1.87	1.29				
	HEMBA1002257	2.5	1.06	1.11	1.47	1.72	1.38	1.46	2.73	1.02				
	HEMBA1002259	3.93	2.57	3.46	3.84	3.35	3.79	1.58	3.6	2.24				

Table 179

	HEMBA1001866	3.9	2.3	1.44	4.16	4.87	4.12	3.87	2.04	2.67				
	HEMBA1001869	9.74	8.73	4.94	27.07	27.58	25.58	12.15	11.95	13.97	**	+	*	+
5	HEMBA1001871	74.25	58.85	43.65	34.31	39.06	32.3	22.21	20.99	22.52			*	-
	HEMBA1001876	3.15	3.01	2.05	6.71	7.01	5.67	24.3	20.84	22.31	**	+	**	+
	HEMBA1001878	8.91	7.59	5.14	7.69	6.34	6.19	2.57	4.4	3.62			*	-
	HEMBA1001879	6.77	3.64	3.77	7.79	7.79	8.38	5.4	7.09	7.12	*	+		
	HEMBA1001884	8.03	4.66	4.9	8.15	7.93	9.25	2.34	3.47	2.61				
10	HEMBA1001886	15.37	8.23	7.45	18.06	17.92	20.6	6.22	8.67	8.91	*	+		
	HEMBA1001888	4.74	2.28	2.28	8.53	6.01	5.71	3.99	3.23	5.19	*	+		
	HEMBA1001890	6.82	5.35	4.39	17.01	13.2	14.58	10.35	9.13	10.28	**	+	**	+
	HEMBA1001896	7.21	3.51	4.27	4.48	4.55	6.32	4.5	3.56	4.29				
	HEMBA1001899	10.27	5.12	6.13	12.84	16.36	13.59	19.93	20.02	20.79	*	+	**	+
15	HEMBA1001904	117.8	90.63	69.63	121.8	145.7	135.1	54.06	69.53	68.48				
	HEMBA1001910	2.98	1.61	1.31	1.77	1.8	2.33	2.01	1.92	2.16				
	HEMBA1001911	24.54	11.64	15.86	17.52	15.24	14.86	10.3	9.59	10.07				
	HEMBA1001912	20.82	8.69	15.18	15.64	15.33	18.75	6.84	9.35	7.93				
	HEMBA1001913	11.57	4.6	5.78	9.2	8.02	9.12	5.36	7.66	8.31				
20	HEMBA1001915	2.07	1.75	1.56	2.72	4.13	3.37	2.79	1.65	1.94	*	+		
	HEMBA1001918	2.07	1.25	1.13	3.95	3.76	3.13	1.5	2.66	1.53	**	+		
	HEMBA1001921	7.05	7.38	3.11	5.25	3.04	7.8	3.53	3.11	2.74				
	HEMBA1001931	0.78	1.98	0.41	1.78	1.48	1.79	0.69	1.82	0.96				
	HEMBA1001939	2.45	1.1	1.29	2.61	2.56	3.15	2.04	3.08	2.2				
25	HEMBA1001940	3.74	2.59	1.93	4.33	6.11	5.9	2.78	3.06	3.22	*	+		
	HEMBA1001942	3.67	2.27	1.69	2.35	3.04	3.41	1.26	2.11	2.03				
	HEMBA1001944	9.44	4.28	2.7	6.72	6.77	6.95	5.78	5.16	5.81				
	HEMBA1001945	2.07	0.91	0.94	1.56	3.05	1.77	1.66	1.79	2.71				
	HEMBA1001950	4.31	3.64	2.4	3.3	1.98	4.19	2.53	3.33	2.77				
30	HEMBA1001951	11.47	5.14	7.18	8.76	8.49	10.31	7.11	7.14	6.62				
	HEMBA1001958	5.93	3.29	3.76	7.31	5.94	5.87	2.95	3.04	4.22				
	HEMBA1001960	5.09	2.29	3.83	2.58	2	3.56	3.69	2.82	3.05				
	HEMBA1001962	0.53	0.49	0.61	0.68	0.72	0.97	-0.01	1.07	0.54				
	HEMBA1001964	1.04	0.26	1.15	2.39	2.99	2.5	0.67	1.12	1.07	**	+		
	HEMBA1001967	5.08	3.46	3.83	6.72	5.35	6.55	3.95	4.57	3.93	*	+		
35	HEMBA1001979	2.59	1.65	1.24	2.97	3.02	3.75	2.54	2.41	2.4	*	+		
	HEMBA1001987	6.47	2.58	3.01	7.96	9.29	7.63	5.55	5.23	5.01	*	+		
	HEMBA1001991	7.79	3.05	3.16	10.3	8.9	8.81	6.21	4.84	5.65	*	+		
	HEMBA1002003	6.67	2.83	3.92	3.54	4.68	6.3	5.41	4.34	5.17				
	HEMBA1002005	4.44	1.76	2.03	5.73	4.88	5.69	3.58	2.87	3.42	*	+		
40	HEMBA1002008	2.92	0.92	1.99	4.42	4.45	4.33	2.3	2.71	2.6	*	+		
	HEMBA1002018	7.24	3.29	3.8	4.79	5.31	4.52	3.14	4.37	3.39				
	HEMBA1002022	0.68	0.34	0.54	1.12	1.17	1.66	0.59	0.97	1.25	*	+		
	HEMBA1002029	147.9	114.2	64.17	209.3	183.3	187.5	83.85	70.94	83.09	*	+		
	HEMBA1002030	3.84	2.17	1.78	2.59	2.01	2.76	1.95	2.52	1.44				
45	HEMBA1002035	4.53	2.83	2.27	3.74	3.23	4.73	2.32	2.93	2.77				
	HEMBA1002037	7.19	3.71	4.11	7.77	6.62	7.18	5.2	4.49	4.12				
	HEMBA1002038	5.05	3.39	2	4.89	4.12	6.29	3.56	4.65	2.86				
	HEMBA1002039	2.43	1.42	2.68	4.62	4.34	5.48	2.31	3.78	2.6	**	+		
	HEMBA1002042	5.07	5.1	4.66	5.37	6.66	7.8	3.75	3.26	4.84				
50	HEMBA1002043	9.02	4.29	4.09	8.45	7.53	9.32	5.8	6.07	6.51				
	HEMBA1002048	3.59	2.88	2.34	3.02	3.12	3.4	3.49	2.47	3.92				
	HEMBA1002049	6.44	2.94	4.68	7.87	9.3	8.66	5.4	4.91	5.09	*	+		
	HEMBA1002053	6.69	4.81	4.26	7.69	7.89	9.03	5.94	6.61	5.76	*	+		
	HEMBA1002055	9.71	8.18	6.93	9.3	5.31	10.84	11.8	6.23	11.57				
55	HEMBA1002056	10.47	4.85	5.55	4.12	3.5	3.57	2.73	3.84	2.21				
	HEMBA1002061	2.87	2.19	2.53	7.31	4.68	4.5	2.4	3.51	2.69	*	+		
	HEMBA1002080	60.84	42.27	48.29	35.05	22.5	22.95	22.84	15.7	24.41	*	-	**	-

Table 178

	HEMBA1001651	21.79	9.98	12.75	16.31	17.89	15.92	12.62	14.58	15.88				
	HEMBA1001655	4.81	3.57	3.37	4.17	7.59	5.82	3.99	4.47	4.7				
5	HEMBA1001658	2.18	2.11	2.13	1.33	1.53	2.6	1.84	1.15	1.86				
	HEMBA1001661	8.45	3.05	2.97	4.66	4.8	5.77	3.88	4.28	4.23				
	HEMBA1001665	5.86	2.62	4.27	4.6	3.94	3.51	4.69	4.17	4.52				
	HEMBA1001670	4.7	2.98	3.53	6.5	7.04	7.21	4.56	5.94	4.89	**	+		
	HEMBA1001672	2.9	1.62	1.17	2.74	2.64	2.91	2.23	3.35	2.84				
10	HEMBA1001673	9.39	3.95	5.37	12.29	9.95	9.16	6.04	3.4	6.06				
	HEMBA1001675	2.77	1.09	1.9	3.14	3.42	1.99	2.07	3.04	2.09				
	HEMBA1001676	66.2	42	41.28	59.83	62.25	61.28	35.33	41.76	48.98				
	HEMBA1001678	23.82	16.82	12.46	26.08	27.44	24.59	15.29	14.2	16.03				
	HEMBA1001680	7.07	3.71	3.69	6.51	7.15	6.71	4.41	4.86	5.34				
15	HEMBA1001681	1.95	0.92	1.52	1.86	1.78	2.38	1.26	2.56	1.49				
	HEMBA1001684	10.32	4.07	5.37	13.29	14.64	14.01	8.6	7.77	8.12	*	+		
	HEMBA1001695	1.84	2.2	0.62	1.62	1.54	2.31	1.72	2.13	0.77				
	HEMBA1001702	3.21	1.66	2.35	4.83	3.35	4.17	3.17	4.1	3.6				
	HEMBA1001709	3.9	1.96	2.65	5.53	4.06	6.56	5.94	7.83	7.54		**	+	
20	HEMBA1001711	2.38	2.81	1.61	5.64	7.85	8.65	3.33	2.8	5.34	**	+		
	HEMBA1001712	2.87	1.69	2.03	2.84	2.47	3.33	3.23	2.26	2.84				
	HEMBA1001714	27.51	15.33	17.22	17.64	16.58	15.17	22.02	17.65	27.85				
	HEMBA1001717	1.6	0.57	0.95	1.72	1.13	1.76	8.51	5.96	6.55		**	+	
	HEMBA1001718	3.34	3.04	3.56	7.23	5.88	7.76	3.79	4.78	3.44	**	+		
25	HEMBA1001723	3.28	1.43	2.31	5.16	4.28	5.3	2.9	4.31	2.84	*	+		
	HEMBA1001731	2.16	1.22	2.13	2.79	1.84	2.37	1.77	2.95	2.23				
	HEMBA1001734	2.33	0.57	2.06	3.71	2.97	2.91	2.16	2.87	2.2				
	HEMBA1001736	8.5	4.87	4.76	7.17	7.6	9.06	7.56	6.14	10.7				
	HEMBA1001741	1.43	1.25	0.91	2.83	2.87	2.84	0.76	1.93	1.43	**	+		
30	HEMBA1001744	1.28	0.91	0.85	1.4	1.01	1.73	0.65	1.88	1.22				
	HEMBA1001745	3.12	1.1	1.48	2.46	2.57	2.63	2.55	3.03	2.51				
	HEMBA1001746	1.85	2.08	1.47	2.46	2.29	3.39	2.8	3.77	3.54		*	+	
	HEMBA1001761	4.88	2.73	3.04	7.7	5.44	7.35	2.96	5.66	4.32	*	+		
	HEMBA1001762	1.84	0.76	1.19	2.52	2.18	2.84	1.18	3.82	1	*	+		
35	HEMBA1001781	3.69	1.25	2.05	4.27	2.77	4.83	2.36	3.3	2.22				
	HEMBA1001784	5.2	3.84	2.76	3.59	2.92	3.32	3.06	2.91	4.28				
	HEMBA1001791	11.2	5.23	3.55	8.42	10	8.96	7.67	7.29	10.19				
	HEMBA1001794	16.08	14.18	10.1	17.79	20.03	18.56	11.08	17.68	19.33	*	+		
	HEMBA1001800	3.13	2.01	2.42	2.99	2.87	3.62	3.26	1.68	2.16				
40	HEMBA1001803	1.53	0.75	0.44	1.21	1.11	1.38	1.41	1.74	1.3				
	HEMBA1001804	13.32	7.17	6.9	11.34	8.08	9.13	6.64	6.79	7.2				
	HEMBA1001808	2.99	2.64	1.45	3.65	1.42	1.93	2.32	1.6	1.79				
	HEMBA1001809	8.19	6.19	4.29	9.5	5.47	7.96	5.87	6.3	5.81				
	HEMBA1001811	22.78	13.64	9.05	9.98	11.16	7.43	11.09	9.04	12.53				
	HEMBA1001815	6.31	3.66	3.82	9.75	9.09	7.86	4.84	3.89	4.99	*	+		
45	HEMBA1001816	2.42	2.34	2.86	3.29	2.61	3.73	2.83	1.8	2.99				
	HEMBA1001819	6.29	5.74	3.76	9.76	8.91	7.84	4.59	4.2	3.42	*	+		
	HEMBA1001820	0.7	1.31	0.35	1.28	0.85	0.69	1	1.34	0.88				
	HEMBA1001822	14.41	5.14	6.88	15.5	12.29	15.38	7.82	5.44	7.89				
	HEMBA1001824	8.95	4.75	6.46	8.72	13.86	15.64	6.51	9.15	7.54				
50	HEMBA1001835	1.68	1.35	0.6	1.34	1.74	3.48	1.4	1.22	1.48				
	HEMBA1001844	7.57	4.41	3.42	10.11	10.97	6.9	6.82	4.22	6.02				
	HEMBA1001847	7.9	4.44	3.77	6.2	5.53	6.41	3.82	3.06	3.92				
	HEMBA1001849	8.79	2.94	4.27	12.39	11.89	10.26	7.27	5.07	7.41	*	+		
	HEMBA1001850	7.06	2.55	2.83	6.48	6.52	7.6	4.57	4.3	4.89				
55	HEMBA1001861	1.79	0.52	1.23	3.32	3.68	4.21	1.63	1.37	1.83	**	+		
	HEMBA1001862	20.07	14.3	16.51	10.49	11.03	16.7	24.43	27.68	20.89		*	+	
	HEMBA1001864	1.89	1.29	0.85	2.99	3	2.74	1.08	1.87	1.04	**	+		

Table 177

	HEMBA1001407	5.43	1.65	2.98	3.95	4.01	3.47	2.95	2.92	2.93			
	HEMBA1001411	2.17	0.69	0.63	2.51	1.83	3.63	1.29	1.35	1.63			
5	HEMBA1001413	5.49	2.49	2.2	4.28	3.2	3.97	3.24	2.49	2.68			
	HEMBA1001414	3.79	2.32	2.38	3.06	1.8	2.44	2.65	3.55	3.21			
	HEMBA1001415	6.49	2.16	2.76	5.46	6.84	6.46	4.32	4.17	5.11			
	HEMBA1001416	6.22	3.74	3.23	8.62	6.54	6.82	5.91	4.3	6.02			
10	HEMBA1001432	5.37	2.98	3.43	7.69	6.86	7.06	3.39	4.18	4.43	*	+	
	HEMBA1001433	4.8	2.47	2.21	6.26	5.3	4.79	3.29	2.49	2.37			
	HEMBA1001435	8.18	4.71	5.41	14.34	11.54	13.2	6.78	6.29	7.26	**	+	
	HEMBA1001442	1.65	1.46	0.73	2.67	3.31	2.57	0.77	1.88	2.03	*	+	
	HEMBA1001446	9.08	2.53	3.23	6.88	6.71	6.42	5.95	6.22	6.04			
15	HEMBA1001450	7.08	5.32	4.43	8.06	5.46	8.96	5.99	5.4	5.68			
	HEMBA1001454	10.16	4.17	5.03	16.08	14.78	15.21	9.95	9.22	10.42	**	+	
	HEMBA1001455	1.25	1.28	0.63	2.33	2.23	1.74	2.53	2.34	2.01	*	+	**
	HEMBA1001459	3.35	1.42	1.26	1.85	2.02	1.94	1.14	1.39	2.31			
	HEMBA1001461	8.81	3.16	4.05	10.82	10.26	6.95	6	5.33	4.95			
	HEMBA1001462	2.66	2.42	2.15	2.1	1.78	2.07	1.34	1.53	2.31			
20	HEMBA1001463	7.17	2.73	3.52	7.24	7.08	8.95	4.33	5.14	4.39			
	HEMBA1001469	7.79	8.03	2.81	8.15	8.71	7.67	5.88	4.2	6.47			
	HEMBA1001473	2.06	0.9	0.31	1.64	1.59	1.3	1.54	1.11	1.32			
	HEMBA1001477	1.25	0.8	0.62	0.91	1.28	0.76	1.34	2.38	1.44			
25	HEMBA1001478	2.09	0.93	1.34	1.5	1.78	0.98	1.62	2.3	1.59			
	HEMBA1001488	12.07	6.47	7.53	8.82	7.12	9.89	6	6.87	5.33			
	HEMBA1001483	4.46	3.27	2.35	2.86	3.34	4.48	1.86	2.27	1.82			
	HEMBA1001490	1.81	1.4	1.03	1.82	1.46	1.52	1.48	2.37	1.32			
	HEMBA1001495	36.22	21.61	21.87	15.42	21.1	17.04	-16.21	19.62	20.73			
	HEMBA1001497	7.26	3.96	4.28	11.8	9.61	9.85	5.21	4.28	5.2	*	+	
30	HEMBA1001510	13.72	5.93	6.56	13.7	15.62	12.58	10.78	9.6	9.58			
	HEMBA1001515	2.6	2	0.87	2.75	3.2	2.93	2.35	3.19	2.52			
	HEMBA1001517	1.89	1.95	1.22	2.95	2.33	2.76	1.72	1.66	2.42	*	+	
	HEMBA1001522	3.61	1.7	1.12	1.99	2.84	1.73	1.04	1.87	1.3			
	HEMBA1001526	5.16	2.43	3.68	6.63	4.1	5.88	3.55	3.16	3.42			
35	HEMBA1001533	8.95	4.93	4.41	7.97	8.75	10.67	4.59	5.06	4.92			
	HEMBA1001547	35.19	25.44	22.4	15.45	14.19	13.27	6.7	4.99	4.47	*	**	-
	HEMBA1001552	8.07	6.24	3.86	9.62	10.94	7.97	8.18	5.74	5.97			
	HEMBA1001553	16.17	10.48	11.7	14.97	19.64	15.26	19.38	22.7	26.62	*	+	
	HEMBA1001557	8.77	5.74	4.35	8.02	8.99	7.7	7.33	5.59	10.39			
40	HEMBA1001563	3.9	1.92	1.89	5.08	3.9	4.71	2.33	3.96	2.78			
	HEMBA1001566	3.98	2.49	2.79	5.22	9.83	5.76	3.59	4.31	4.01			
	HEMBA1001569	8.8	4.36	5.19	13.14	14.49	14.76	6.66	7.84	10.58	**	+	
	HEMBA1001570	10.01	5.49	7.22	16.18	15.76	21.41	6.88	8.18	7.08	*	+	
	HEMBA1001579	14.95	9.44	8.88	11.45	10.82	11.3	6.85	6.64	9.61			
45	HEMBA1001581	6.6	2.62	2.74	9.65	8.35	7.34	4.2	4.87	7.29	*	+	
	HEMBA1001582	1.39	1.89	0.99	1.46	1.52	1.21	1.87	1.43	1.14			
	HEMBA1001585	3.5	1.76	2.06	4.04	4.61	4.34	2.06	2.32	2.78	*	+	
	HEMBA1001589	5.07	3.16	2.15	3.41	3.1	3.21	3.05	2.93	3.94			
	HEMBA1001595	13.49	4.3	6.8	10.71	10.28	9.89	6.83	8.2	8.67			
50	HEMBA1001604	5.72	2.28	3.75	6.52	7.03	5.34	2.89	3.22	3.58			
	HEMBA1001606	8.03	3.96	3.18	8.15	6.4	9.15	2.3	4.25	3.65			
	HEMBA1001615	46.6	32.92	22.49	33.05	34.32	33.44	126.5	104.9	149.7	**	+	
	HEMBA1001620	14.48	8.32	7.64	17.62	16.71	15.28	9.29	10.91	11.09	*	+	
	HEMBA1001621	9.93	5.63	3.68	7.55	6.93	6.59	5	4.37	6.21			
	HEMBA1001635	5.73	3.82	2.42	4.14	4.05	5.67	2.93	3.94	3.69			
55	HEMBA1001636	4.39	1.44	3.08	3.97	3.02	3.88	3.71	3.49	4.55			
	HEMBA1001640	3.49	0.97	1.46	3.57	2.02	3.07	2.4	3.06	2.05			
	HEMBA1001647	6.4	2.49	4.47	3.2	5.99	5.63	2.28	4.03	4.16			

Table 176

HEMBA1001221	2.19	1.18	0.78	2.28	2.36	2.53	2.4	3.63	1.51				
HEMBA1001225	1.21	1.77	1.22	2.62	2.13	1.37	0.82	1.74	2.82				
HEMBA1001226	13.52	10.49	8.9	18.36	20	19.62	7.7	10.44	7.45	**	+		
HEMBA1001228	13.05	5.12	4.29	9.55	8.22	7.69	6.04	7.48	7.86				
HEMBA1001229	12.71	9.28	6.69	8.25	7.48	7.38	10.2	8.81	12.42				
HEMBA1001235	4.86	4.97	4.74	7.89	8.06	6.71	5.12	7.06	11.33	**	+		
HEMBA1001238	5.14	3.54	3.32	7.04	6.92	8.57	3.98	4.55	5.25	*	+		
HEMBA1001242	9.9	9.56	8.33	13.88	6.68	13.26	5.82	6.16	5.11		**	-	
HEMBA1001247	4.46	1.61	1.9	3.57	3.49	3.72	3.78	3.48	3.42				
HEMBA1001253	5.27	3.3	2.61	4.73	4.85	2.62	2.61	2.92	2.88				
HEMBA1001257	3.88	2.26	2.32	3.08	5.15	4.69	1.41	2.58	1.9				
HEMBA1001261	30.79	16.66	18.37	18.07	18.08	21.82	20.19	23.46	27.67				
HEMBA1001262	2.76	4.04	1.52	6.54	5.42	3.57	2.84	3.16	4.61				
HEMBA1001265	5.3	6.7	4.27	9.23	8.19	10.09	4.34	5.27	5.82	*	+		
HEMBA1001266	7.76	6.62	6.38	9.89	9.6	8.87	6.28	5.38	7.65	**	+		
HEMBA1001269	37.26	20.56	22.9	18.88	18.77	19.35	8.45	11.29	14.06		*	-	
HEMBA1001272	1.9	1.41	1.17	1.81	2.19	2.98	1.62	1.83	1.14				
HEMBA1001279	7.18	4.55	5.66	6.03	6.98	6.47	3.39	5.47	3.9				
HEMBA1001281	5.42	5.55	6.33	11.93	16.02	13.78	5.82	4.84	7.89	**	+		
HEMBA1001286	25.93	14.58	10.17	19.52	21.27	19.41	15.05	12.01	17.84				
HEMBA1001289	4.9	3.9	2.72	4.42	4.59	5.54	4.24	2.99	5.3				
HEMBA1001291	12.14	5.79	5.07	8.25	5.62	6.51	5.37	5.12	8.98				
HEMBA1001294	3.24	2.44	2.03	4.94	4.48	4.82	2.73	2.45	3.08	**	+		
HEMBA1001296	3.68	1.37	1.28	2.91	2.24	3.02	2.56	2.34	2.65				
HEMBA1001297	5.4	4.74	4.72	5.79	6.42	4.8	3.21	2.6	2.27		**	-	
HEMBA1001299	6.03	3.81	4.28	7.69	11.74	10.72	5.99	5.39	5.03	*	+		
HEMBA1001302	6.53	3.1	5.55	4.99	5.75	7.13	4.2	5.14	4.56				
HEMBA1001303	3.57	2.21	0.92	2.41	4.91	3.42	1.52	2.66	2.14				
HEMBA1001306	22.18	12.36	12.24	18.89	23.21	22.17	16.22	12.41	17.9				
HEMBA1001308	11.41	6.87	7.33	12.58	12.35	13.73	8.36	8.24	9.57	*	+		
HEMBA1001310	7.91	5.67	6.18	9.02	7.1	8.4	7.65	6.89	8.59				
HEMBA1001312	6.83	4.78	4.59	4.91	5.69	6.9	6.83	6.24	6.66				
HEMBA1001319	0.37	0.17	0.45	0.79	0.92	1.12	0.66	2.44	0.75	**	+		
HEMBA1001322	7.21	5.19	6.74	8.06	10.08	9.08	6.21	7.42	7.75	*	+		
HEMBA1001323	4.23	3.25	2.82	10.32	10.14	7.03	8.56	8.82	9.24	**	+	**	+
HEMBA1001326	5.74	3.25	2.25	3.17	5.59	5.42	5.13	3.49	5.64				
HEMBA1001327	2.36	2.51	1.03	2	2.41	3.09	2.74	2.46	3.87				
HEMBA1001330	5.82	5.46	4.35	11.86	14.54	13.29	6.08	7.36	9.3	**	+		
HEMBA1001348	3.13	2.19	2.78	4.2	2.23	2.88	1.71	2.63	2.74				
HEMBA1001350	12.36	10.68	7.51	15.66	13.69	14.52	11.25	9.44	10.45	*	+		
HEMBA1001351	8.18	6.48	5.91	13	14.47	12.39	10.67	8.35	8.14	**	+		
HEMBA1001352	7.26	6.11	6.06	7.73	6.7	6.17	5.26	7.29	6.09				
HEMBA1001353	31.3	26.87	27.53	25.75	22.23	20.82	12.94	17.74	19.72	*	-	**	-
HEMBA1001358	34.05	17.05	14.31	20.81	35.28	26	9.32	9.14	12.44				
HEMBA1001361	1.82	1.14	2.1	2.53	3.2	3.65	3	2.26	2.92	*	+	*	+
HEMBA1001364	1.53	0.54	0.65	1.45	1.91	1.58	2.49	1.92	1.51				
HEMBA1001375	3.85	2.39	2.36	2.46	4.27	4.44	3.43	3.13	3.87				
HEMBA1001377	8.53	6.83	5.73	14.04	14.14	13.21	6.15	8.25	6.71	**	+		
HEMBA1001383	2.54	1.25	1.73	3.25	2.57	3.3	1.34	2.99	1.96				
HEMBA1001387	4.07	1.84	3.25	5.31	4.33	4.27	3.43	4.91	3.24				
HEMBA1001398	4.68	4.67	4.87	6.78	7.58	7.44	4.52	4.77	5.63	**	+		
HEMBA1001390	7.44	5.12	5.37	11.6	11.73	11.98	10.29	9.6	11.06	**	+	**	+
HEMBA1001391	1.33	1.22	1.11	3.9	2.02	2.65	2.4	1.36	1.99	*	+		
HEMBA1001398	5.47	2.84	3.19	6.23	6.58	7.24	3.41	4.42	5.15	*	+		
HEMBA1001405	5.26	1.42	2.09	2.65	2.11	2.81	3.1	2.12	3.28				
HEMBA1001406	3.16	2.03	2.11	4.74	3.74	4.54	2.55	3.04	2.7	*	+		

Table 175

	HEMBA1000979	5.49	2.18	2.97	6.7	3.77	4.39	3.48	5.27	4.03			
5	HEMBA1000981	9.63	9.63	8.99	5.49	6.85	5.43	3.2	5.8	4.89	**	-	**
	HEMBA1000983	6.43	3.92	2.91	5.46	7.35	6.51	4.3	3.18	4.68			
	HEMBA1000985	1.63	1.32	0.83	1.53	0.96	1.83	1.43	0.82	1.18			
	HEMBA1000986	8.66	3.3	4.89	7.79	10.67	12.32	6.59	5.63	7.52			
	HEMBA1000991	3.99	3.51	3.27	7.03	8.03	8.59	3.11	5.46	4.41	**	+	
10	HEMBA1001007	6.98	3.16	4.1	4.53	6.32	6.25	5.08	5.14	4.03			
	HEMBA1001008	3.18	2.08	1.67	6.05	4.43	4.59	2.99	3.85	3.36	*	+	
	HEMBA1001009	3.19	2.06	1.89	3	2.73	3.35	2.83	4.13	2.55			
	HEMBA1001014	5.39	3.12	5.74	9.86	11.08	12.45	4.65	7.98	7.55	**	+	
	HEMBA1001017	7.4	4.83	4.74	5.73	6.28	5.4	4.08	4.41	5.88			
15	HEMBA1001019	2.85	2.29	1.26	2.91	2.72	2.07	1.51	2.11	2.14			
	HEMBA1001020	3.1	1.76	1.25	4.02	4.91	3.89	2.56	2.42	2.65	*	+	
	HEMBA1001021	5.67	3.26	3.56	5.27	3.84	4.59	5.11	3.82	6.55			
	HEMBA1001022	4.52	3.09	3.23	5.25	4.72	3.27	2.64	3.83	3.89			
	HEMBA1001024	1.94	0.42	0.87	1.28	1.11	2.19	1.54	1.4	1.01			
	HEMBA1001026	1.87	1.27	0.7	1.76	2.89	2.28	1.38	1.06	1.68			
20	HEMBA1001043	2.16	1.91	1.95	3.51	4.01	3.96	1.57	1.82	0.63	**	+	
	HEMBA1001051	12.22	4.76	5.28	19.03	15.88	16.82	10.42	7.53	10.73	*	+	
	HEMBA1001052	1.62	0.97	1.98	2.53	4.21	2.8	2.24	1.49	2.61			
	HEMBA1001059	6.89	2.24	2.49	4.96	3.77	4.85	4.31	4.18	4.43			
	HEMBA1001060	7.98	3.88	4.72	10.32	9.35	8.51	6.1	5.55	6.56	*	+	
25	HEMBA1001064	5.36	3.84	3.22	6.43	5.68	4.77	2.55	3.39	3.71			
	HEMBA1001071	1.62	1.41	0.32	16	17.18	11.61	12.79	12.04	12.64	**	+	**
	HEMBA1001077	4.45	3.8	1.96	11.6	9.35	8.57	3.08	5.61	3.95	**	+	
	HEMBA1001078	14.1	8.18	8.99	5.43	6.25	7.02	4.32	6.96	5.16			
	HEMBA1001080	5.79	3.95	2.49	3.69	5.23	5.89	5.35	4.03	3.93			
30	HEMBA1001084	5.31	2.86	2.62	7.71	7.07	6.47	5.73	4.4	5.39	*	+	
	HEMBA1001085	13.38	7.46	10.01	19.29	18.48	14.18	11.36	11.18	10.99	*	+	
	HEMBA1001088	5.8	4.05	4.96	5.45	4.2	4.92	5.6	5.06	6.59			
	HEMBA1001093	2.01	1.13	0.59	2.57	2.37	1.64	1.63	2.12	1.53			
	HEMBA1001094	0.9	1.06	0.61	2.27	2.81	2.04	1.48	1.38	2.02	**	+	*
35	HEMBA1001099	2.64	3.87	2.39	4.48	2.58	3.18	1.73	2.49	1.54			
	HEMBA1001104	4.32	2.56	3.02	5.08	3.19	2.29	3.64	4.68	2.66			
	HEMBA1001109	15.93	10.15	10.15	27.48	26.01	22.62	15.71	11.93	11.35	**	+	
	HEMBA1001114	8.6	5.78	5.64	9.84	9.77	10.41	14.65	11.13	18.58	*	+	*
	HEMBA1001121	2.07	1.57	0.99	2.33	3.89	3.11	2.34	1.82	1.7	*	+	
40	HEMBA1001122	2.51	5.06	1.5	14.85	12.94	9.66	6.46	7.06	7.13	**	+	*
	HEMBA1001123	10.26	5.27	4.03	8.74	8.81	11.74	6.7	7.3	6.19			
	HEMBA1001133	4.14	2.91	3.18	3.04	2.73	4.12	2.58	3.25	4.04			
	HEMBA1001137	9.39	4	4.74	6.72	8.14	6.94	3.8	6.14	4.6			
	HEMBA1001140	6.82	5.7	6.11	10.25	12.69	12.18	4.71	6.45	5.99	**	+	
45	HEMBA1001144	14.92	3.84	7.57	18.27	23.75	20.85	12.2	8.33	12.65	*	+	
	HEMBA1001145	28.51	33.95	19.22	28.92	30.82	30.17	44.7	41.59	36.72		*	+
	HEMBA1001158	5.04	3.15	2.61	5.99	3.8	6.16	5.34	3.86	4.7			
	HEMBA1001172	5.81	3.09	2.82	8.57	8.02	8.53	5.72	4.06	4.92	*	+	
	HEMBA1001174	2.3	2.42	1	1.59	2.22	1.73	1.99	2.72	1.46			
	HEMBA1001175	4.94	2.83	3.63	9.64	8.74	8.9	6.25	5.58	6.09	**	+	*
50	HEMBA1001182	15.48	8.24	12.75	16.98	16.95	14.52	6.34	9.16	8.55			
	HEMBA1001184	1.37	1.11	1.17	2.46	2.2	1.94	1.7	1.64	1.09	**	+	
	HEMBA1001192	1.14	1.3	0.71	1.4	1.96	2.74	1.17	1.75	1.29			
	HEMBA1001196	9.67	6.82	7.53	12.04	9.93	10.61	8.76	5.56	6.62			
	HEMBA1001197	26.77	18.72	19.29	43.93	39.7	25.47	17.6	13.26	17.88			
55	HEMBA1001208	4.45	2.51	2.06	2.43	4.11	3.2	2.93	2.71	2.12			
	HEMBA1001213	4.18	2.48	2.29	4.96	5.22	3.94	3.68	3.27	5.1			
	HEMBA1001214	28.24	15.89	17.42	11.21	10.37	12.48	8.74	7.69	7.51		*	-

Table 174

	HEMBA1000719	4.82	2.97	2.79	3.61	4.58	3.67	3.75	2.77	3.67				
	HEMBA1000722	2.03	0.86	1.42	1.98	2.82	1.59	1.34	3.92	2.07				
5	HEMBA1000726	10.3	9.3	7.72	23.56	26.89	19.83	12.69	13.58	11.3	**	+	*	+
	HEMBA1000727	6.04	3.96	3.25	8.14	10.98	7.59	6.32	6.82	2.98	*	+		
	HEMBA1000732	3.01	2.28	1.42	2.14	1.87	1.92	2.98	2.21	2.48				
	HEMBA1000736	4.72	2.16	2	3.64	1.97	1.99	2.73	2.2	2.64				
	HEMBA1000743	0.32	1.05	0.53	1.51	2.41	0.98	0.72	1.22	1.24				
10	HEMBA1000745	1.74	1.73	1.32	1.18	1.69	2.12	1.96	2.53	1.18				
	HEMBA1000747	4.19	1.78	1.08	3.03	2.21	1.78	1.85	3.32	2.09				
	HEMBA1000748	2.17	1.28	2.24	2.2	3.52	2.79	1.6	2.38	1.72				
	HEMBA1000749	4.95	3.09	2.17	6.45	8.33	7.14	3.25	4.29	3.58	*	+		
	HEMBA1000752	4.81	3.6	2.79	5.03	6.01	4.99	3.34	3.06	3.28				
15	HEMBA1000753	9.91	6.17	6.18	9.28	11.1	8.29	5.77	5.12	5.5				
	HEMBA1000757	7.1	7.74	5.44	11.01	14.04	12.37	5.58	4.46	4.75	**	+		
	HEMBA1000760	16.78	13.36	13.64	8.72	12.16	6.16	8.22	7.22	7.97		**	-	
	HEMBA1000769	7.05	2.51	3.23	9	8.67	9.72	4.24	4.83	3.98	*	+		
	HEMBA1000773	1.32	0.68	0.25	0.36	1.46	1.1	0.81	1.64	0.68				
20	HEMBA1000774	8	3.27	7.05	12.39	12.55	13.92	7.51	8.12	7.46	*	+		
	HEMBA1000780	2.14	1.77	0.74	2.61	2.17	1.75	1.28	2.13	1.21				
	HEMBA1000783	1.08	1.96	1.07	2.21	1.08	2.2	1.9	1.74	1.44				
	HEMBA1000791	3.14	3.15	3.13	6.58	7.55	5.76	3.73	3.72	6.22	**	+		
	HEMBA1000793	9.3	4	3.98	5.49	6.95	5.86	5.38	4.76	5.7				
25	HEMBA1000802	3.76	2.25	1.22	2.43	3.6	2.62	0.88	2.18	1.88				
	HEMBA1000813	9.81	3.16	4.27	6.99	7.53	7.12	3.67	6.02	6.65				
	HEMBA1000817	2.66	1.43	0.92	2.74	3.08	2.72	1.26	2.52	1.67				
	HEMBA1000822	0.99	1.09	0.85	1.62	3.22	2.71	1.22	1.82	0.71	*	+		
	HEMBA1000827	7.7	6.4	3.84	6.01	6.66	6.53	3.91	3.03	4.64				
30	HEMBA1000833	5.1	2.66	2.23	8.93	7.69	7.93	7.69	5.86	6.86	**	+	*	+
	HEMBA1000835	5.71	3.29	3.29	5.75	3.34	4.85	2.51	3.39	3.41				
	HEMBA1000843	6.36	5.57	5.21	6.61	9.85	9.29	4.9	5.64	10.02				
	HEMBA1000851	4.2	1.79	2.1	3.58	3.85	2.86	2.91	1.96	2.78				
	HEMBA1000852	5.4	3.22	2.28	5.81	4.07	5.82	2.77	3.99	3.71				
35	HEMBA1000867	1.61	2.47	1.06	2.17	3.19	2.37	0.68	2.24	0.83				
	HEMBA1000869	1.82	1.11	0.72	0.98	2.58	1.99	0.79	2.22	0.83				
	HEMBA1000870	6.82	3.33	3.67	6.25	6.67	4.52	3.47	4.37	5.69				
	HEMBA1000872	4.12	2.25	3.08	4.7	5.64	4.68	3.33	3.29	4.33	*	+		
	HEMBA1000875	1.77	1.41	1.93	5.81	7.31	5.85	7.19	6.68	8.14	**	+	**	+
	HEMBA1000876	5.86	4.79	3.07	7.1	7.28	6.57	4.55	4.52	6.23	*	+		
40	HEMBA1000907	2.12	2.01	0.66	2.54	2.27	2.12	2.3	1.55	1.2				
	HEMBA1000908	4.73	8.03	3.2	3.97	8	4.77	4.32	3.17	3.88				
	HEMBA1000910	4.06	2.39	3.23	5.88	8	5.6	3.31	3.17	3.05	*	+		
	HEMBA1000918	3.62	1.79	2.38	3.54	2.97	3.56	2.53	2.34	2.18				
	HEMBA1000919	6.44	3.37	2.05	4.74	4.83	4.38	3.75	4.79	3.18				
45	HEMBA1000934	8.7	4.01	3.95	4.96	5.39	5.6	4.1	3.51	4.76				
	HEMBA1000935	2.09	1.32	1.09	2.05	2.33	2	1.1	2.66	1.72				
	HEMBA1000940	4.94	2.14	2.53	3.07	4.88	4.53	2.3	2.63	3.13				
	HEMBA1000942	6.3	3.89	2.49	7.11	9.28	8.54	5.01	6.29	5.19	*	+		
	HEMBA1000943	1.76	1.55	1.49	3.18	2.76	2.23	1.98	2.58	2.05	*	+	*	+
50	HEMBA1000946	8.15	7.73	6	5.16	5.33	4.88	2.92	2.97	2.91	*	-	**	-
	HEMBA1000960	9.59	5.75	7.08	15.65	18.02	18.53	8.6	9.11	11.03	**	+		
	HEMBA1000962	6.47	2.77	4.32	4.75	7	7.13	5.16	3.51	7.23				
	HEMBA1000968	7	1.7	1.54	3	4.17	3.31	2.23	2.96	3.14				
	HEMBA1000971	5.14	1.71	2.36	4.85	4.32	4.5	3.47	4.15	3.21				
55	HEMBA1000972	3.69	1.13	1.73	5.98	4.9	5.9	2.76	4.55	2.35	*	+		
	HEMBA1000974	1.6	0.93	0.68	2.29	1.66	2.44	2.01	3.61	2.23	*	+		
	HEMBA1000975	3.28	2	1.5	5.97	3.13	2.57	2.25	4.14	2.32				

Table 173

	HEMBA1000481	20.13	11.47	12.73	18.55	18.55	15.53	7.84	7.33	12.91				
	HEMBA1000488	7.66	4.44	4.62	7.86	6.19	6.89	3.5	5.38	6.42				
5	HEMBA1000490	4.18	2.68	1.34	3.95	5.37	3.63	2.12	2.88	4.31				
	HEMBA1000491	7.15	3.43	2.52	5.5	6.82	6.64	4.25	3.29	3.33				
	HEMBA1000498	10.26	6.11	4.98	10.58	18.06	18.44	9.53	6.44	8.57	*	+		
	HEMBA1000501	10.31	9.16	7.08	7.41	5.02	8.46	4.06	4.46	3.72		**	-	
	HEMBA1000504	0.29	1.06	0.88	2.55	1.79	2.74	3.2	4.91	2.54	*	+	*	+
10	HEMBA1000505	4	3.11	2.61	4.34	3.87	4.06	3.11	3.95	3.94				
	HEMBA1000507	8.99	4.59	6.64	9.35	10.47	8.65	5.55	8.59	7.24				
	HEMBA1000508	8.59	6.68	6.07	11.49	13.9	16.57	7.32	8.75	9.79	*	+		
	HEMBA1000518	2.98	1.78	1.55	2.04	2.31	1.71	2.15	1.54	1.87				
	HEMBA1000519	13.74	9.63	6.41	18.15	26.1	23.45	14.61	12.39	16.75	*	+		
15	HEMBA1000520	0.74	1.54	1.42	0.53	4.99	5.32	0.3	3.24	3.21				
	HEMBA1000523	2.58	1.73	1.85	2.49	2.81	3.42	2.38	3.31	2.63				
	HEMBA1000531	5.39	5.46	3.11	3.93	6.67	3.26	3.72	3.54	2.94				
	HEMBA1000534	0.79	3.21	2.91	1.73	9.74	6.64	0.85	6.6	3.17				
	HEMBA1000538	-0.07	2.6	2.6	0.69	6.28	5.42	0.12	7.11	5.18				
20	HEMBA1000540	3.94	2.64	3.3	8.03	7.49	8.11	2.04	3.68	2.54	**	+		
	HEMBA1000542	5.67	3.4	2.44	3.85	3.5	5.44	3.98	3.82	4.97				
	HEMBA1000545	2.41	1.53	0.38	4.15	3.69	3.21	1.98	2.16	2.09	*	+		
	HEMBA1000547	1.74	1.59	1.68	5.72	8.77	7.03	3.43	3.74	3.3	**	+	**	+
	HEMBA1000551	9.65	6.1	8.03	14.99	17.46	18.61	8.56	8.89	9.19	**	+		
25	HEMBA1000555	5.3	2	2.07	3.79	6.18	4.25	2.7	2.98	2.37				
	HEMBA1000557	4.48	2.92	3.57	7.15	7.8	8.32	4.31	6.14	5.01	**	+		
	HEMBA1000561	3.7	1.44	1.77	4.14	3.06	3.15	3.47	4.41	2.34				
	HEMBA1000563	1.24	0.37	0.85	2.27	1.82	2.27	0.66	2.98	0.86	*	+		
	HEMBA1000567	3.87	1.04	1.51	8.01	8.19	8.67	2.66	3.73	4	**	+		
30	HEMBA1000568	3.88	2.11	2.05	5.69	5.23	5.4	1.77	2.82	3.91	*	+		
	HEMBA1000569	4.97	2.5	2.71	6.85	4.01	5.8	3.46	3.51	4.29				
	HEMBA1000575	13.92	7.22	8.43	20.52	24.59	18.68	11.63	11.79	11.04	*	+		
	HEMBA1000588	1.28	0.91	1.2	2.91	2.49	2.9	1.78	2.48	2.62	**	+	*	+
	HEMBA1000590	3.14	1.5	1.84	3.09	1.65	1.71	1.44	1.82	1.81				
	HEMBA1000591	6.68	3.59	4.87	8.78	6.73	9.08	5.54	5.94	6.27				
35	HEMBA1000592	1.77	1	1.66	2.61	3.4	2.25	1.98	2.18	1.99	*	+		
	HEMBA1000594	3.25	0.68	1.19	1.74	3.07	2.12	1.39	1.15	1.72				
	HEMBA1000604	5.99	4.47	2.05	8.88	9.05	6.96	6.29	5.91	6.23	*	+		
	HEMBA1000607	4.99	3.1	3.35	6.44	6.82	5.81	3.43	4.28	4.42	*	+		
	HEMBA1000608	0.99	1.94	0.42	3.85	2.15	1.46	2.61	2.1	3.4				
40	HEMBA1000622	2.66	1.16	0.99	4.04	3.67	4.04	2.76	3.15	3.26	*	+		
	HEMBA1000634	28.82	15.23	16.08	35.62	36.93	32.2	24.35	21.77	26.76	*	+		
	HEMBA1000636	10.44	4.41	5.46	7.42	7.72	8.03	6.42	4.97	5.75				
	HEMBA1000637	5.28	3.33	4.09	4.63	6.26	5.53	4.14	4.87	4.43				
	HEMBA1000655	7.39	4.24	2.84	8.57	9.07	9.85	5.75	6.56	6.78	*	+		
45	HEMBA1000657	7.14	3.75	3.78	6.89	5.66	6.19	7.09	4.53	7.57				
	HEMBA1000662	2.8	1.64	1.1	1.89	1.7	1.33	1.86	1.9	1.81				
	HEMBA1000664	2.6	2.45	0.17	3.74	3.57	2.7	2.86	2.52	2.77				
	HEMBA1000671	3.69	2.81	2.74	7.05	5.05	5.15	3.14	2.82	3.51	*	+		
	HEMBA1000673	5.96	2.79	3.34	9.32	7.79	7.67	4.47	3.8	5.32	*	+		
50	HEMBA1000675	2.45	2.8	0.77	6.63	4.04	4.43	3.65	3.8	3.87	*	+	*	+
	HEMBA1000678	7.03	5.09	6.34	10.12	8.74	9.2	2.93	5.72	5.28	*	+		
	HEMBA1000682	5.22	2.07	2.75	12.42	15.95	13.04	14.17	11.88	14.92	**	+	**	+
	HEMBA1000686	5.1	3.46	2.35	5.21	4.74	3.32	3.54	2.67	2.25				
	HEMBA1000702	9.79	6.15	6.42	10.8	11.22	8.35	8.93	8.45	8.8				
55	HEMBA1000705	1.79	1.26	0.4	2.12	2.25	1.15	1.75	1.57	2.17				
	HEMBA1000713	5.65	3.58	2.89	6.69	5.36	6.21	7.06	6.72	4.64				
	HEMBA1000718	4.7	2.67	2.33	5.7	6	5.76	3.69	3.85	2.59	*	+		

Table 172

	HEMBA1000231	5.77	4.79	1.39	5.73	5.06	5.05	2.57	4.02	3.83				
	HEMBA1000237	10.5	9.41	7.28	13.8	14.47	14.03	8.59	13.21	9.08	**	+		
5	HEMBA1000243	4.4	2.18	1.57	4.11	5.36	4.88	3.72	3.39	3.4				
	HEMBA1000244	11.09	6.03	5.16	9.66	7.12	6.2	9.02	6.06	9.63				
	HEMBA1000251	2.83	2.17	1.02	2.88	4.48	2.64	1.69	2.92	2.44				
	HEMBA1000254	5.6	3.06	2.15	6.61	5.66	5.33	3.44	3.21	4.84				
	HEMBA1000264	3.12	2.38	1.29	3	2.42	2.07	2.39	1.18	3.05				
10	HEMBA1000269	3.15	2.65	1.66	4.09	3.3	1.89	1.88	1.49	1.6				
	HEMBA1000275	10.1	8.27	6.59	12.65	12.4	13.32	7.47	7.72	5.65	*	+		
	HEMBA1000280	2.4	1.67	1.88	3.2	3.34	2.25	0.92	2.83	1.47				
	HEMBA1000282	4.3	2.15	1.99	8.2	7.71	7.54	4.05	3.59	4.68	**	+		
	HEMBA1000287	6.5	5	3.8	6.66	6.95	7.33	6.19	6.14	4.66				
15	HEMBA1000288	4.22	5.47	1.6	5.44	4.7	5.08	3.8	2.7	3.03				
	HEMBA1000290	2.44	1.68	1.41	3.3	2.07	2.24	2.46	1.37	1.82				
	HEMBA1000296	4.58	3.23	3.04	3.88	4.57	3.87	2.97	3.13	3.49				
	HEMBA1000300	7.18	7.47	4.77	15.63	12.41	11.86	8.05	9.96	6.36	**	+		
	HEMBA1000302	2.87	1.87	1.42	2.86	2.56	2.8	1.34	2.59	1.57				
20	HEMBA1000303	12.63	6.43	5.95	8.6	9.24	8.52	6.4	8.51	7.91				
	HEMBA1000304	5.94	4.85	2.91	8.58	10.98	8.79	6.22	5.73	5.36	*	+		
	HEMBA1000307	3.35	2.83	1.79	7.52	6.27	5.03	5.57	4.79	3.97	*	+	*	+
	HEMBA1000312	7.59	5.13	7.25	13.4	9.35	10.01	7.66	6.43	8.25	*	+		
	HEMBA1000318	4.73	3.46	2.76	7.07	6.34	4.78	4.52	5.17	4.75				
25	HEMBA1000327	4.9	14.95	2.36	5.69	8.99	5.72	3.18	5.4	3.63				
	HEMBA1000333	2.68	1.29	0.21	2.59	1.6	1.38	2.24	1.33	1.95				
	HEMBA1000338	7.1	5.92	3.55	10.42	12.67	10.27	5.82	7.1	5.05	*	+		
	HEMBA1000343	4	2.99	2.01	2.63	3.79	2.89	1.22	2.1	1.84				
	HEMBA1000349	3.15	2.72	2.94	1.9	3.38	2.84	1.58	1.8	2.44		*		
30	HEMBA1000351	12.26	4.06	4.63	9.54	11.2	9.66	5.66	5.25	4.95				
	HEMBA1000355	5.83	4.02	3.82	5.03	5.09	4.09	3.9	3.77	4.2				
	HEMBA1000356	8.5	4.16	3.88	9.66	6	7.29	7.01	5.23	5.35				
	HEMBA1000357	6.36	2.11	3.61	7.55	7.35	8.12	3.8	3.56	3.53	*	+		
	HEMBA1000366	2.01	1.56	0.82	2.54	1.86	2.67	1.26	2.04	1.96				
35	HEMBA1000369	7.61	3.99	4.13	5.06	4.64	5.24	3.29	3.78	3.59				
	HEMBA1000370	1.94	1.23	1.23	3.73	3.06	3.01	1.19	2.46	1.97	**	+		
	HEMBA1000376	5.48	4.4	4.48	8.19	9.77	8.68	4.81	5.75	4.74	**	+		
	HEMBA1000387	6.72	4.8	4.24	12.88	11.31	8.93	7.04	6.86	7.9	*	+		
	HEMBA1000389	6.41	4.31	3.18	5.44	5.19	3.87	3.91	4.16	5.13				
	HEMBA1000390	2.89	3.46	2.42	2.82	2.5	3.02	2.55	2.1	2.56				
40	HEMBA1000392	1.66	1.01	0.96	2.76	2.9	2.64	1.17	2.08	1.89	**	+		
	HEMBA1000396	2.67	1.46	1.17	3.48	2.29	1.9	2.07	2.04	2.6				
	HEMBA1000411	2.73	2.11	2	2.49	2.83	1.98	1.3	2.58	1.84				
	HEMBA1000418	2.29	2.59	1.6	3.21	4.57	2.67	2.11	3.04	2.45				
	HEMBA1000422	5.88	3.82	2.78	5.71	5.46	6.46	2.91	5	3.36				
45	HEMBA1000428	2.98	1.47	1	5.92	5.67	4.87	3.36	3.17	3.89	**	+		
	HEMBA1000434	0.46	1.18	0.48	1.51	2.2	1.01	1.46	1.36	1.4		*	+	
	HEMBA1000442	1.91	1.74	2.18	1.99	2.71	2.66	1.77	2.2	1.7				
	HEMBA1000443	5.28	4.21	2.77	4.95	5.35	7.43	4.57	4.71	4				
	HEMBA1000446	15.47	8.43	7.47	8.86	8.46	9.56	8.97	8.38	10.15				
50	HEMBA1000456	7.87	3.87	5.62	12.88	11.2	12.65	6.87	8.86	10.32	**	+		
	HEMBA1000459	3.86	2.75	1.81	4.89	5.61	4.96	2.29	3.47	3.74	*	+		
	HEMBA1000460	2.95	1.91	1.24	1.69	3.46	2.84	3.05	2.46	5.23				
	HEMBA1000462	17.16	10.03	4.79	13.14	13.57	10.69	11.49	13.69	11.75				
	HEMBA1000464	1.23	1.41	0.6	1.41	1.89	0.9	1.32	1.26	0.96				
	HEMBA1000468	1.87	1.63	0.67	3.5	1.61	1.75	2.85	2.43	2.2				
55	HEMBA1000469	4.36	2.95	2.67	7.93	8.36	9.97	5.39	4.1	4.79	**	+		
	HEMBA1000477	6.04	2.58	2.34	5.17	5.61	5.34	6	5.59	6.01				

Table 171

	BRAWH1000060	2.9	2.93	1.8	3.46	3.35	2.78	2.07	3.22	2.32				
	BRAWH1000075	2.06	1.78	1.17	2.08	2.99	2.28	1.92	2.13	2.14				
5	BRAWH1000081	4.56	1.87	2.1	2.75	2.22	2.25	1.42	2.46	1.85				
	BRAWH1000084	26.93	16.26	13.57	23.37	33.3	27.71	19.86	27.26	24.74				
	BRAWH1000095	11.47	5.88	3.86	6.15	6.04	6.04	6.03	4.2	5.03				
	BRAWH1000096	7.17	5.2	3.04	5.76	6.13	4.73	6.35	5.93	7.43				
	BRAWH1000097	7.61	5.42	4.3	8.36	9.37	10.77	5.92	6.56	7.12 *	+			
10	BRAWH1000100	2.35	1.26	1.29	3.27	4.09	3.18	3.47	3.17	3.82 *	+	+		
	BRAWH1000101	15.93	5.73	7.58	15.78	16.69	15.33	10.38	7.98	10.75				
	BRAWH1000104	1.83	1.99	1.25	3.05	2.31	2.64	0.9	2.83	2.28 *	+			
	BRAWH1000107	5.24	3.06	2.55	3.69	4.48	3.14	2.51	6.62	2.54				
	BRAWH1000110	37.02	23.89	17.95	52.01	48.45	48.78	25.83	19.88	30.82 *	+			
15	BRAWH1000111	13.78	8.87	6.05	12.15	10.84	10.06	10.64	8.06	9.74				
	BRAWH1000135	11.51	6.6	6.16	7.34	6.27	6.18	7.86	5.16	9.04				
	BRAWH1000190	5.57	3.61	3.06	4.88	4.05	4.63	4.28	3.62	5.01				
	HEMBA1000005	2.17	2.36	2.39	3.59	3.26	3.09	2.51	1.69	3.76 **	+			
	HEMBA1000006	4.88	4.08	3.07	5.64	5.07	4.69	3.89	4.34	3.69				
20	HEMBA1000012	7.67	9.97	9.83	7.99	7.06	6.98	3.55	5.22	3.46		**		
	HEMBA1000020	27.06	14.56	16.3	24.94	23.65	29.76	15.51	14.38	17.35				
	HEMBA1000030	7.2	6.04	4.37	4.93	6.66	4.71	4.8	4.96	7.17				
	HEMBA1000034	5.42	3.03	3.13	3.92	5.81	5.55	2.45	2.65	5.55				
	HEMBA1000042	10.53	5.34	5.29	12.34	15.71	15.33	6.74	5.14	8.81 *	+			
25	HEMBA1000045	3.35	1.45	2	3.11	2.27	3.63	2.78	2.42	2.82				
	HEMBA1000046	4.44	3.21	3.62	6.34	8.01	11.1	5.61	5.39	6.03 *	+	**	+	
	HEMBA1000047	3.38	2.86	1.36	3.03	2.25	2.95	2.29	1.9	1.25				
	HEMBA1000048	6.35	3.98	4.34	16.75	14.72	14.62	7.09	8.13	7.75 **	+	+	+	
	HEMBA1000050	1.73	0.67	0.56	1.86	1.47	1.56	1.52	2.71	1.56				
30	HEMBA1000053	2.66	1.5	1.58	2.81	3.5	3.13	2.37	1.92	3.37 *	+			
	HEMBA1000060	4.78	3.18	2.77	4.56	4.67	4.59	3.9	4.27	4.27				
	HEMBA1000072	71.82	55.54	44.63	47.17	62.62	63.43	25.66	24.24	32.66		*		
	HEMBA1000073	2.41	1.46	1.48	2.36	2.35	2.6	1.84	2.72	2.72				
	HEMBA1000076	10.02	11.17	8.35	27.94	21.02	20.27	16.4	9.49	15.31 **	+			
35	HEMBA1000084	3.64	2.86	3.72	4.85	4.96	4.11	5.09	5.98	4.83 *	+	+	+	
	HEMBA1000087	3.12	2.56	2.1	4.7	3.46	2.58	2.59	4.09	3.28				
	HEMBA1000088	1.57	0.55	0.65	1.47	0.74	0.92	1.69	2.19	2.78		*	+	
	HEMBA1000091	7.82	3.65	3.58	5.14	4.68	5.32	5.87	2.69	5.02				
	HEMBA1000111	3.34	2.33	2.42	4.87	5.39	5.9	3.66	3.37	3.36 **	+			
40	HEMBA1000121	3.69	2.19	1.8	4.54	7.02	6.59	3.95	3.3	4.32 *	+			
	HEMBA1000128	4.07	1.73	1.88	3.07	3.61	4.19	4.82	5.85	5.45		*	+	
	HEMBA1000129	4.83	2.28	2.77	2.81	3.65	3.39	2.57	2.73	3.94				
	HEMBA1000141	2.71	2.09	1.62	4.16	2.77	4.01	2.77	3.67	1.66 *	+			
	HEMBA1000146	2.9	1.3	1.8	2.65	2.28	1.73	1.61	3.65	1.85				
45	HEMBA1000150	26.65	13.33	17.02	31.39	35.61	38.63	19.78	16.66	26.75 *	+			
	HEMBA1000154	36.53	16.72	17.93	24.12	23.55	16.21	9	9.29	13.92				
	HEMBA1000156	12.63	7.55	7.2	12.13	11.18	10.85	5.44	6.27	10.52				
	HEMBA1000158	14.24	5.92	4.83	15.57	17.46	14.26	10.9	12.16	12.71				
	HEMBA1000168	10.07	5.72	5.58	8.47	10.06	8.07	7.36	7.05	5.56				
	HEMBA1000180	3.67	1.14	1.34	3.4	2.55	2.88	1.78	2.08	2.49				
50	HEMBA1000185	9.44	4.05	4.26	11.55	10.93	10.36	7.42	5.5	5.94 *	+			
	HEMBA1000188	2.86	1.61	0.93	2.94	2.35	3.1	1.57	1.58	1.71				
	HEMBA1000193	1.27	0.58	0.24	1.37	0.89	0.82	0.26	0.53	0.45				
	HEMBA1000194	11.09	4.55	5.41	17.15	17.6	13.81	11.08	8.03	17.29 *	+			
	HEMBA1000201	3.51	1.9	1.75	4.07	2.62	2.46	2.06	2.69	2.83				
55	HEMBA1000213	2.2	0.91	0.97	1.85	2.66	1.89	1.72	1.64	1.67				
	HEMBA1000216	4.38	3.53	3.49	7.1	6.02	3.1	3.46	3.84	4.14				
	HEMBA1000227	6.93	1.95	2.95	5.37	3.71	3.99	3.84	2.55	3.65				

Clone	NT2			NT2 RA			NT2 RA INHIB			test	+	test	+
	exp.1	exp.2	exp.3	exp.1	exp.2	exp.3	exp.1	exp.2	exp.3	N/R	-	N/I	-
GAPDH(Cr1)	3.53	1.08	0.98	2.92	2.49	2.8	1.76	2.59	1.52				
β actin(Cr2)	155.4	118	99.68	148.5	110.7	101.3	114.7	105.8	151.1				
ADRGL1000005	4.01	2.03	1.55	4.05	3.65	3.6	2.27	2.93	4.24				
ADRGL1000007	11.08	5.73	7.92	15.42	10.6	13.87	8.99	8.17	9.15				
ADRGL1000009	1.11	0.72	1.04	1.66	1.89	1.03	1.22	1.62	1.58		*		+
ADRGL1000011	4.27	2.7	2.85	4.32	4.35	3.38	2.76	3.27	3.06				
ADRGL1000027	1.83	0.38	0.56	0.97	0.62	0.99	0.92	1.33	1.5				
ADRGL1000058	3.65	2.58	1.37	2.92	3.36	2.75	2.25	3.51	2.7				
ADRGL1000069	3.25	1.85	3.28	1.86	2.53	2.85	2.01	2.89	2.7				
ADRGL1000077	13.48	10.41	6.71	19.62	17.92	22.59	11.6	16.66	19.34	*	+		
ADRGL1000092	5.73	2.8	4.51	7.31	5.01	4.83	3.24	6.16	7.22				
ADRGL1000099	5.64	3.42	2.08	5.59	3.73	4.24	3.98	3.98	4.06				
ADRGL1000136	9.97	3.52	4.19	5.77	4.73	5.86	6.61	5.16	5.49				
ADRGL1000147	23.09	13.85	11.7	14.77	14.96	14.89	17.7	13.3	19.47				
ADRGL1000159	6.11	2.22	3.37	5.24	2.88	4.15	2.76	2.93	3.59				
ADRGL1000160	7.16	3.48	4.19	5.94	4.59	3.41	3.95	4.67	4.25				
ADRGL1000171	4.84	2.99	3.23	3.52	4.19	4.37	2.55	3.88	3.45				
ADRGL1000181	5.1	3.65	2.6	3.16	4.06	2.97	2.64	3.06	3.44				
BGGI11000015	13.95	6.83	6.72	9.61	9.19	10.24	9.94	10.66	10.13				
BGGI11000016	15.49	5.92	7.09	11.88	11.38	8.72	11.82	10.98	10.51				
BGGI11000017	7.89	2.99	3.25	4.94	4.94	4.93	3.55	4.27	3.52				
BGGI11000022	8.77	5.14	5.91	7.12	7.05	4.54	5.71	5.59	5.9				
BGGI11000031	4.71	2.16	2.74	4.09	3.29	3.96	4.02	3.67	2.33				
BGGI11000042	6.37	5.24	3.74	5.63	6.22	4.36	4.66	5.2	4.04				
BGGI11000046	19.01	12.57	9.23	12.39	15.7	12.37	8.8	10.92	9.17				
BNGH41000020	859	910.1	603	164	319.2	267.4	638.2	771.6	845.4	**	-		
BNGH41000025	5.35	2.06	2.09	2.76	2.76	3.77	4.23	2.01	3.06				
BNGH41000026	16.2	7.69	7.05	9.34	11.37	9.66	10.13	7.16	10.71				
BNGH41000027	2.31	2.18	2.5	2.9	3.01	2.82	3.68	3.48	4.21	**	+	**	+
BNGH41000035	14.57	8.83	9.36	10.92	9.55	14.75	15.02	15.18	12.2				
BNGH41000037	10.56	7.46	6.2	8.16	9.21	6.42	3.37	5.45	4.98				
BNGH41000042	77.1	50.85	58.45	47.64	53.39	62.67	28.12	35.48	23.44		*		-
BNGH41000048	3.5	2.19	1.91	4.28	2.87	2.4	1.63	3.01	1.78				
BNGH41000056	2.57	2.01	1	1.91	2.63	2.15	1.41	2.4	1.79				
BNGH41000087	9.84	5.84	5.53	12.49	10.24	10.25	11.74	9.68	8.53				
BNGH41000091	3.37	2.59	1.21	3.29	3.01	1.55	2.95	2.57	2.13				
BNGH41000157	10.63	5.64	6.15	8.53	9.05	7.74	6.38	6.68	5.75				
BNGH41000169	3.77	4.34	3.82	4.9	3.48	3.32	3.4	4.16	4.19				
BNGH41000181	2.47	1.59	1.84	2.93	2.1	1.8	1.7	2.66	1.59				
BNGH41000198	8.13	4.64	3.79	5.48	4.35	5.59	4.3	4.15	4.35				
BNGH41000219	9.61	3.92	4.87	4.17	5.29	5.45	5.24	7.12	7.13				
BNGH41000239	19.61	13.28	8.68	10.86	11.27	9.36	7.9	9.5	10.85				
BNGH41000237	10.9	5.47	6.45	6.65	6.97	7.79	6.36	6.25	5.44				
BNGH41000238	4.58	7	3.45	5.91	4.68	4.34	4.33	5.44	4.22				
BNGH41000243	13.85	8.69	8.48	10.19	9.71	8.97	8.23	4.87	5.54				
BNGH41000270	5.83	2.62	2.35	2.3	3.05	3.44	2.59	3.49	1.3				
BRAWH1000004	4.19	2.83	2.48	5.04	3.15	3.26	1.44	3.45	2.05				
BRAWH1000018	4.85	1.95	2.29	7.47	8.8	8.85	8.68	6.61	7.96	**	+	*	+
BRAWH1000021	6.52	5.06	5.87	5.09	6.94	6.44	2.89	6.23	4.28				
BRAWH1000027	11.64	8.86	7.19	8.24	10.39	11.51	5.58	7.13	8.24				
BRAWH1000029	9.58	5.15	3.52	6.01	6.72	6	5.08	5.12	5.84				
BRAWH1000040	4.6	1.89	2.14	2.92	2.71	2.7	2.92	2.5	3.01				
BRAWH1000050	11.48	4.95	5.19	9.74	7.25	8.62	8.25	8.09	8.93				
BRAWH1000051	8.18	3.93	3.19	6.15	5.72	6.02	5.01	4.25	4.44				

	Y79AA1002355	48.88	42.39	40.68	0.87	0.83
	Y79AA1002361	87.11	88.66	76.9	1.02	0.88
5	Y79AA1002365	38.75	24.26	20.53	1	1
	Y79AA1002373	43.96	55.06	28.34	1.25	0.91
	Y79AA1002376	3080.7	3824.05	4481.1	1.24	1.45
	Y79AA1002378	73.33	93.61	68.22	1.28	0.93
10	Y79AA1002381	248.36	288.51	304.13	1.16	1.22
	Y79AA1002388	118.82	135.82	129.37	1.14	1.09
	Y79AA1002399	36.12	30.1	32.87	1	1
15	Y79AA1002407	57.84	42.82	52.54	0.74	0.91
	Y79AA1002413	78.77	81.36	87.31	1.03	1.11
	Y79AA1002416	34.3	30.2	51.99	1	1.3
	Y79AA1002429	67.91	69.81	80.19	1.03	1.18
20	Y79AA1002431	24.66	21.16	23.98	1	1
	Y79AA1002433	27.12	18.11	23.63	1	1
	Y79AA1002445	78.66	54.58	73.75	0.69	0.94
	Y79AA1002461	29.04	24.84	32	1	1
25	Y79AA1002466	882.69	904.65	782.53	1.02	0.89
	Y79AA1002471	53.74	51.26	68.91	0.95	1.28
	Y79AA1002472	121.95	127.4	127.11	1.04	1.04
	Y79AA1002474	53.33	40.85	47.18	0.77	0.88
30	Y79AA1002482	103.36	111.11	116.07	1.07	1.12
	Y79AA1002487	30.92	25.8	32.51	1	1
	Y79AA1002490	101.4	90.92	90.54	0.9	0.89
	Y79AA1002493	107.88	125.54	105.75	1.16	0.98
35	ZRV6C1006278	46.63	30.08	32.23	0.86	0.86

Table 170

Expression of each cDNA in undifferentiated NT2 cells, in NT2 cells cultured in the presence of retinoic acid, or in NT2 cells that were cultured in the presence of retinoic acid and then further cultured in the presence of cell-division inhibitor added (This table also contains clones without description in Examples)

In the table, NT2, NT2_RA, and NT2_RA_INHIB represent untreated NT2 cells, retinoic acid-treated NT2 cells, and retinoic acid/inhibitor-treated NT2 cells, respectively. The assay was performed in triplicate (n=3), and each result was shown in the column of exp.1, exp.2, or exp.3. In addition, "t-test N/R" and "t-test N/I" represent results of test for significance of difference between the untreated cells and the retinoic acid-treated cells, and between the untreated cells and the retinoic acid/inhibitor-treated cells, respectively. The results of the test are shown in the columns of *:p<0.05 and **:p<0.01.

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	Y79AA1002067	69.44	85.04	88.86	1.22	1.28
	Y79AA1002069	22.45	14.55	11.72	1	1
	Y79AA1002070	290.21	261.7	420.51	0.9	1.45
5	Y79AA1002074	4784.8	4322.02	3500.57	0.9	0.73
	Y79AA1002076	23.92	22.89	30.01	1	1
	Y79AA1002083	32.64	25.47	18.78	1	1
	Y79AA1002084	56.93	47.45	58.61	0.83	1.03
10	Y79AA1002086	29.98	30.26	23.7	1	1
	Y79AA1002087	339.84	473.89	305.27	1.39	0.9
	Y79AA1002089	63.57	68.96	66.81	1.08	1.05
	Y79AA1002093	45.69	38.45	36.91	0.88	0.88
15	Y79AA1002101	19.61	22.84	19.2	1	1
	Y79AA1002103	37.61	37.75	35.74	1	1
	Y79AA1002115	48.3	60.52	68.39	1.25	1.42
	Y79AA1002121	50.52	36.73	32.42	0.79	0.79
20	Y79AA1002125	81.65	86.76	79.2	1.06	0.97
	Y79AA1002129	33.63	31.52	18.59	1	1
	Y79AA1002131	20.65	14.34	20.31	1	1
	Y79AA1002139	31.44	25.87	27.48	1	1
25	Y79AA1002144	138.16	90.77	145.45	0.66	1.05
	Y79AA1002177	53.67	57.36	47.56	1.07	0.89
	Y79AA1002183	168.93	154.43	165.5	0.91	0.98
	Y79AA1002202	111.2	133.56	100.48	1.2	0.9
30	Y79AA1002204	35.34	28.87	17.92	1	1
	Y79AA1002206	28.51	22.35	18.61	1	1
	Y79AA1002208	36.92	36.47	44.08	1	1.1
	Y79AA1002209	48.12	45.04	49.5	0.94	1.03
	Y79AA1002210	29.11	36.94	29.41	1	1
35	Y79AA1002211	57.36	63.16	52.25	1.1	0.91
	Y79AA1002213	133.51	136.73	123.74	1.02	0.93
	Y79AA1002215	139.31	112.01	119.56	0.8	0.86
	Y79AA1002220	69.79	59.74	58.27	0.86	0.83
40	Y79AA1002226	91.29	74.46	87.97	0.82	0.96
	Y79AA1002229	34.29	24.76	28.27	1	1
	Y79AA1002234	34.52	56.44	51.95	1.41	1.3
	Y79AA1002235	73.58	67.73	74.78	0.92	1.02
45	Y79AA1002246	50.76	51.58	35.73	1.02	0.79
	Y79AA1002258	63.15	82.64	90.15	1.31	1.43
	Y79AA1002279	122.18	148.88	110.84	1.22	0.91
	Y79AA1002292	61.42	72.22	67.83	1.18	1.1
	Y79AA1002298	39.81	43.55	30.57	1.09	1
50	Y79AA1002307	32.77	30.65	23.26	1	1
	Y79AA1002309	18.06	24.11	13.63	1	1
	Y79AA1002311	53.31	57.19	84.46	1.07	1.58
	Y79AA1002334	35.05	38.07	31.31	1	1
55	Y79AA1002351	42.13	61.09	58.56	1.45	1.39

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	Y79AA1001719	56.65	50.42	51.8	0.89	0.91
	Y79AA1001727	63.97	82.15	75.03	1.28	1.17
	Y79AA1001750	201.52	244.27	213.32	1.21	1.06
5	Y79AA1001760	1525.1	1826.08	1836.66	1.2	1.2
	Y79AA1001777	27.57	22.01	20.75	1	1
	Y79AA1001781	14.2	10.34	11.67	1	1
	Y79AA1001787	34.67	32.77	31.46	1	1
10	Y79AA1001793	490.12	500.64	415.02	1.02	0.85
	Y79AA1001795	20.82	15.64	16.66	1	1
	Y79AA1001799	63.3	90.19	82.19	1.42	1.3
	Y79AA1001800	206.49	248.75	358.98	1.2	1.74
15	Y79AA1001801	59.81	46	44.71	0.77	0.75
	Y79AA1001803	32.78	24.73	24.59	1	1
	Y79AA1001805	93.54	111.29	88.98	1.19	0.95
	Y79AA1001807	156.64	104.4	119.1	0.67	0.76
20	Y79AA1001827	54.55	36.34	49.28	0.73	0.9
	Y79AA1001846	90.02	81.03	88.49	0.9	0.98
	Y79AA1001848	46.02	35.28	38.62	0.87	0.87
	Y79AA1001853	54.99	60.14	72.48	1.09	1.32
25	Y79AA1001863	55.19	71.44	82.39	1.29	1.49
	Y79AA1001866	73	77.51	62.79	1.06	0.86
	Y79AA1001874	12.88	7.85	8.7	1	1
	Y79AA1001875	92.9	99.69	99.18	1.07	1.07
	Y79AA1001907	1093.8	755.57	1038.49	0.69	0.95
30	Y79AA1001908	28.59	18.2	21.42	1	1
	Y79AA1001923	40.94	40.51	35.36	0.99	0.98
	Y79AA1001927	54.84	54.43	48.17	0.99	0.88
	Y79AA1001930	25.74	40.86	42.58	1.02	1.06
35	Y79AA1001932	97.69	98.39	84.63	1.01	0.87
	Y79AA1001933	57.63	46.41	44	0.81	0.76
	Y79AA1001942	37.83	42.2	33.88	1.06	1
	Y79AA1001963	342.06	324.07	294.47	0.95	0.86
40	Y79AA1001968	133.87	148.65	174	1.11	1.3
	Y79AA1001983	31.53	29.07	29.92	1	1
	Y79AA1002000	38.81	38.99	47.02	1	1.18
	Y79AA1002004	37.99	102.36	131.4	2.56	3.29
45	Y79AA1002008	90.25	65.44	54.66	0.73	0.61
	Y79AA1002012	127.69	126.92	98.41	0.99	0.77
	Y79AA1002017	12.63	3.46	2.78	1	1
	Y79AA1002022	130.63	151.97	106.29	1.16	0.81
	Y79AA1002027	31.61	32.98	29.25	1	1
50	Y79AA1002050	45.8	38.26	51.32	0.87	1.12
	Y79AA1002058	1897.1	1240.87	921.95	0.65	0.49
	Y79AA1002060	63.24	95.77	121.69	1.51	1.92
	Y79AA1002062	138.95	173.15	116.65	1.25	0.84
55	Y79AA1002065	241.79	213.59	256.71	0.88	1.06

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	Y79AA1001299	84.16	104.65	104.16	1.24	1.24
	Y79AA1001312	29.58	27.93	24.52	1	1
	Y79AA1001319	50.23	74.51	92.58	1.48	1.84
5	Y79AA1001323	29.99	42.34	39.19	1.06	1
	Y79AA1001328	54.08	55.7	41.63	1.03	0.77
	Y79AA1001343	4712.6	5708.31	5950.25	1.21	1.26
	Y79AA1001351	30.34	22.22	24.29	1	1
10	Y79AA1001364	57.23	72.81	54.68	1.27	0.96
	Y79AA1001367	33.46	28.07	29.13	1	1
	Y79AA1001384	16.59	11.98	21.5	1	1
	Y79AA1001391	22.75	18.1	18.89	1	1
15	Y79AA1001394	107.41	106.15	85.41	0.99	0.8
	Y79AA1001402	88.44	106.63	80.2	1.21	0.91
	Y79AA1001410	31.55	32.32	27.63	1	1
	Y79AA1001414	89.66	99.17	102.16	1.11	1.14
20	Y79AA1001426	26.19	30.96	29.42	1	1
	Y79AA1001427	944.35	1175.39	818.75	1.24	0.87
	Y79AA1001430	62.12	77.68	66.7	1.25	1.07
	Y79AA1001439	106.67	110.26	142.44	1.03	1.34
	Y79AA1001485	46.51	42.89	31.32	0.92	0.86
25	Y79AA1001493	35.54	24.61	23.88	1	1
	Y79AA1001511	57.74	65.35	93.44	1.13	1.62
	Y79AA1001523	37.34	66.39	30.87	1.66	1
	Y79AA1001530	106.49	109.86	93.6	1.03	0.88
30	Y79AA1001532	99.42	111.62	100.16	1.12	1.01
	Y79AA1001533	64.49	53.61	44.36	0.83	0.69
	Y79AA1001541	23.65	33.4	31.44	1	1
	Y79AA1001548	180.15	182.1	165.86	1.01	0.92
35	Y79AA1001555	57.85	54.82	39.03	0.95	0.69
	Y79AA1001562	159.81	149.99	152.18	0.94	0.95
	Y79AA1001581	54.55	60.53	50.36	1.11	0.92
	Y79AA1001585	96.52	98.24	79.13	1.02	0.82
40	Y79AA1001592	62.47	74.66	67.17	1.2	1.08
	Y79AA1001594	6.58	7.46	4.36	1	1
	Y79AA1001603	1371.8	1354.64	1311.51	0.99	0.96
	Y79AA1001613	41.9	47.86	41.96	1.14	1
	Y79AA1001630	21.43	17.4	21.48	1	1
45	Y79AA1001647	76.41	83.21	65.45	1.09	0.86
	Y79AA1001664	87	149.39	85.61	1.72	0.98
	Y79AA1001665	46.95	45.27	52.61	0.96	1.12
	Y79AA1001679	226.19	250.68	232.31	1.11	1.03
50	Y79AA1001692	36.23	50.52	46.28	1.26	1.16
	Y79AA1001696	14.49	20.08	15.55	1	1
	Y79AA1001705	28.44	37.49	24.51	1	1
	Y79AA1001711	93.7	115.28	83.78	1.23	0.89
55	Y79AA1001717	21.82	24.44	17.74	1	1

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	Y79AA1000989	119.43	125.67	133.77	1.05	1.12
	Y79AA1000991	1240.5	994.5	880.71	0.8	0.71
	Y79AA1001013	1435.4	1460.4	1293.41	1.02	0.9
5	Y79AA1001014	59.34	57.2	61.99	0.96	1.04
	Y79AA1001019	50.26	49.6	55.51	0.99	1.1
	Y79AA1001020	77.34	83.75	108.42	1.08	1.4
	Y79AA1001023	30.63	37.84	28.21	1	1
10	Y79AA1001030	54.37	72.86	67.84	1.34	1.25
	Y79AA1001035	36.18	52.22	60.57	1.31	1.51
	Y79AA1001041	34.02	26.39	39.75	1	1
	Y79AA1001043	104.7	132.62	114.65	1.27	1.1
15	Y79AA1001048	36.81	40.02	39.21	1	1
	Y79AA1001056	83.24	70.9	78.16	0.85	0.94
	Y79AA1001061	109.03	110.8	115.01	1.02	1.05
	Y79AA1001062	29.05	40.37	33.85	1.01	1
20	Y79AA1001068	155.79	171.69	179.73	1.1	1.15
	Y79AA1001073	63.42	240.36	260.57	3.79	4.11
	Y79AA1001077	141.08	122.58	123.69	0.87	0.88
	Y79AA1001078	89.83	99.86	113.12	1.11	1.26
25	Y79AA1001081	52.76	53.5	48.66	1.01	0.92
	Y79AA1001088	430.32	406.89	460.33	0.95	1.07
	Y79AA1001089	92.85	116.61	115.46	1.26	1.24
	Y79AA1001090	67.22	69.72	59.84	1.04	0.89
	Y79AA1001105	122.65	110.5	72.39	0.9	0.59
30	Y79AA1001142	62.73	127.62	148.35	2.03	2.36
	Y79AA1001145	99.57	109.62	105.59	1.1	1.06
	Y79AA1001162	61.53	55.57	54.52	0.9	0.89
	Y79AA1001167	22.2	27.76	22.05	1	1
35	Y79AA1001176	22.58	13.4	14.3	1	1
	Y79AA1001177	33.47	34.18	31.87	1	1
	Y79AA1001179	172.36	288.96	209.97	1.68	1.22
	Y79AA1001185	4.57	59.88	4.92	1.5	1
40	Y79AA1001201	75.33	82.93	99.95	1.1	1.33
	Y79AA1001205	37.54	42.63	48.27	1.07	1.21
	Y79AA1001211	97.72	107.77	98.62	1.1	1.01
	Y79AA1001212	95.32	102.86	78.99	1.08	0.83
45	Y79AA1001216	650.81	671.69	747.64	1.03	1.15
	Y79AA1001228	81.54	86.06	78.83	1.06	0.97
	Y79AA1001233	24.91	19.52	21.84	1	1
	Y79AA1001236	41.43	68.09	69.24	1.64	1.67
50	Y79AA1001239	66.08	63.52	79.5	0.96	1.2
	Y79AA1001240	40.45	44.69	30.33	1.1	0.99
	Y79AA1001255	209.03	243.4	327.16	1.16	1.57
	Y79AA1001264	77.47	114.96	155.28	1.48	2
	Y79AA1001272	94.67	108.27	133.39	1.14	1.41
55	Y79AA1001281	18.21	17.06	16.63	1	1

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	Y79AA1000697	242.38	296.34	355.35	1.22	1.47
	Y79AA1000700	43.92	66.94	58.24	1.52	1.33
	Y79AA1000702	139.07	146.21	285.74	1.05	2.05
5	Y79AA1000704	18.66	25.29	15.37	1	1
	Y79AA1000705	44.31	44.51	31.94	1	0.9
	Y79AA1000717	75.03	98.74	86.2	1.32	1.15
	Y79AA1000722	41.86	34.12	41.61	0.96	0.99
10	Y79AA1000724	72.99	67.48	66.4	0.92	0.91
	Y79AA1000726	33.66	36.24	29.59	1	1
	Y79AA1000734	44.87	42.21	66.46	0.94	1.48
	Y79AA1000748	31.18	25.23	28.06	1	1
15	Y79AA1000750	122.85	138.64	121.01	1.13	0.99
	Y79AA1000752	32.8	31.76	33.14	1	1
	Y79AA1000774	66.23	53.85	62.7	0.81	0.95
	Y79AA1000776	26.59	22.27	24.72	1	1
20	Y79AA1000777	129.38	122.4	131.92	0.95	1.02
	Y79AA1000778	51.22	36.45	32.48	0.78	0.78
	Y79AA1000782	63.92	63.79	87.96	1	1.38
	Y79AA1000784	52.79	64.69	85.36	1.23	1.62
25	Y79AA1000794	23.36	22.73	19.03	1	1
	Y79AA1000800	29.75	32.72	33.2	1	1
	Y79AA1000802	30.35	26.74	20.73	1	1
	Y79AA1000805	19.03	22.52	16.55	1	1
30	Y79AA1000814	106.9	114.58	89.95	1.07	0.84
	Y79AA1000823	158.18	147.37	123.66	0.93	0.78
	Y79AA1000824	36.55	29.44	33.27	1	1
	Y79AA1000827	72.11	68.69	52.85	0.95	0.73
	Y79AA1000831	100.55	112.39	119.5	1.12	1.19
35	Y79AA1000833	2837.1	2936.23	2467.7	1.03	0.87
	Y79AA1000850	56.14	49.87	45.58	0.89	0.81
	Y79AA1000856	86.17	85.24	87.41	0.99	1.01
	Y79AA1000862	21.4	15.44	14.93	1	1
40	Y79AA1000876	126.25	128.25	138.28	1.02	1.1
	Y79AA1000888	391.22	560.15	327.4	1.43	0.84
	Y79AA1000902	91.22	102.61	97.33	1.12	1.07
	Y79AA1000935	63.3	75.01	65.76	1.18	1.04
45	Y79AA1000959	59.37	76.73	60.41	1.29	1.02
	Y79AA1000962	36.15	37.57	34.42	1	1
	Y79AA1000963	96.8	192.62	232.02	1.99	2.4
	Y79AA1000966	519	651.66	477.36	1.26	0.92
50	Y79AA1000967	19.89	292.47	251.96	7.31	6.3
	Y79AA1000968	80.03	76.94	75.11	0.96	0.94
	Y79AA1000969	37.01	40.02	31.68	1	1
	Y79AA1000976	14.88	17.75	13.35	1	1
	Y79AA1000978	51.09	73.65	56.87	1.44	1.11
55	Y79AA1000985	236.63	273.81	258.11	1.16	1.09

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	Y79AA1000307	18. 53	20. 75	18	1	1
	Y79AA1000313	49. 32	51. 4	49. 22	1. 04	1
5	Y79AA1000314	75. 77	84. 06	101. 41	1. 11	1. 34
	Y79AA1000328	29. 85	29. 66	36. 26	1	1
	Y79AA1000334	23. 99	28. 51	35. 56	1	1
	Y79AA1000342	327. 09	366. 24	329. 71	1. 12	1. 01
	Y79AA1000346	81. 6	47. 85	70. 26	0. 59	0. 86
10	Y79AA1000347	133. 77	102. 63	126. 89	0. 77	0. 95
	Y79AA1000349	108. 67	107. 41	141. 16	0. 99	1. 3
	Y79AA1000355	162. 38	124. 35	158. 8	0. 77	0. 98
	Y79AA1000368	300. 14	326. 89	275. 96	1. 09	0. 92
15	Y79AA1000388	264. 81	244. 65	255. 31	0. 92	0. 96
	Y79AA1000392	29. 82	42. 96	63. 94	1. 07	1. 6
	Y79AA1000405	76. 52	90. 11	72. 49	1. 18	0. 95
	Y79AA1000410	344. 28	441. 37	334. 83	1. 28	0. 97
20	Y79AA1000420	51. 5	55. 34	53. 93	1. 07	1. 05
	Y79AA1000423	99. 42	111. 57	110. 84	1. 12	1. 11
	Y79AA1000426	48. 92	57. 41	47. 33	1. 17	0. 97
	Y79AA1000432	31. 27	26. 61	25. 92	1	1
25	Y79AA1000453	79. 54	259. 3	206. 07	3. 26	2. 59
	Y79AA1000465	17. 15	42	49. 86	1. 05	1. 25
	Y79AA1000469	168. 58	190. 42	171. 96	1. 13	1. 02
	Y79AA1000480	43. 91	47. 71	38. 1	1. 09	0. 91
	Y79AA1000502	92. 82	78. 1	86. 82	0. 84	0. 94
30	Y79AA1000521	41. 82	34. 78	40. 15	0. 96	0. 96
	Y79AA1000534	82. 06	125. 38	113. 59	1. 53	1. 38
	Y79AA1000538	184. 83	261. 28	173. 89	1. 41	0. 94
	Y79AA1000539	354. 26	344. 61	222. 22	0. 97	0. 63
35	Y79AA1000540	26. 42	33. 12	31. 6	1	1
	Y79AA1000560	1601. 9	1282. 77	1345. 84	0. 8	0. 84
	Y79AA1000574	24. 19	29. 5	35. 63	1	1
	Y79AA1000584	48. 15	39. 35	38. 38	0. 83	0. 83
40	Y79AA1000589	2778. 1	2470. 28	1588. 55	0. 89	0. 57
	Y79AA1000598	29. 48	26. 04	21. 58	1	1
	Y79AA1000600	114. 94	209. 4	237. 16	1. 82	2. 06
	Y79AA1000609	31. 84	21. 28	28. 15	1	1
45	Y79AA1000618	63. 61	66. 01	76. 25	1. 04	1. 2
	Y79AA1000627	62. 01	61. 57	49. 97	0. 99	0. 81
	Y79AA1000636	92. 84	80. 65	118. 18	0. 87	1. 27
	Y79AA1000649	96. 75	123. 64	122. 08	1. 28	1. 26
50	Y79AA1000656	1766. 0	1687. 12	1253. 96	0. 96	0. 71
	Y79AA1000673	27. 77	24. 97	21. 2	1	1
	Y79AA1000674	1171. 6	1373. 94	651. 44	1. 17	0. 56
	Y79AA1000678	36. 94	44. 09	34. 34	1. 1	1
	Y79AA1000682	1983. 1	1699. 68	1530. 39	0. 86	0. 77
55	Y79AA1000683	49. 85	42. 48	43. 56	0. 85	0. 87

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	VESEN2000430	82.14	123.46	143.79	1.5	1.75
	VESEN2000448	13.1	15.75	14.63	1	1
5	VESEN2000449	77.75	116.78	106.38	1.5	1.37
	VESEN2000456	16.97	16.07	16.5	1	1
	VESEN2000562	141.49	130.91	105.14	0.93	0.74
	VESEN2000573	8.91	8.61	6.79	1	1
	VESEN2000604	13.88	12.29	11.35	1	1
10	VESEN2000614	226.89	314.06	318.98	1.38	1.41
	VESEN2000638	17.11	18.86	13.45	1	1
	VESEN2000641	30.44	33.47	35.1	1	1
	VESEN2000645	114.84	103.19	111.61	0.9	0.97
15	Y79AA1000013	23.65	28.36	17.86	1	1
	Y79AA1000030	54.3	54.62	46.92	1.01	0.86
	Y79AA1000033	111.64	118.12	102.03	1.06	0.91
	Y79AA1000037	88.44	96.39	66.63	1.09	0.75
20	Y79AA1000041	30.18	31.51	34.51	1	1
	Y79AA1000059	55.17	55.82	52.18	1.01	0.95
	Y79AA1000065	270.08	316.25	285.13	1.17	1.06
	Y79AA1000081	261.42	374.34	508.29	1.43	1.94
25	Y79AA1000127	107.56	107.37	68.62	1	0.64
	Y79AA1000130	127.02	147.17	116.76	1.16	0.92
	Y79AA1000131	6111.2	5656.63	5788.88	0.93	0.95
	Y79AA1000134	52.82	69.93	53.91	1.32	1.02
	Y79AA1000143	117.86	127.82	125.02	1.08	1.06
30	Y79AA1000144	74.35	76.59	74.72	1.03	1
	Y79AA1000150	2545.9	2718.65	2054.54	1.07	0.81
	Y79AA1000153	5666.0	5284.5	5846.98	0.93	1.03
	Y79AA1000166	55.33	54.66	50.83	0.99	0.92
35	Y79AA1000179	91.15	94.27	64.88	1.03	0.71
	Y79AA1000181	46.14	50.95	39.78	1.1	0.87
	Y79AA1000202	588.9	770.83	565.25	1.31	0.96
	Y79AA1000207	188.44	235.03	155.99	1.25	0.83
40	Y79AA1000214	266.68	298.79	319.39	1.12	1.2
	Y79AA1000222	86.44	101.81	106.59	1.18	1.23
	Y79AA1000226	82.03	161.73	169.02	1.97	2.06
	Y79AA1000227	48.01	65.96	45.99	1.37	0.96
45	Y79AA1000230	31.14	25.61	22.03	1	1
	Y79AA1000231	118.41	136.39	99.08	1.15	0.84
	Y79AA1000239	151.15	168.81	146.2	1.12	0.97
	Y79AA1000258	29.42	44.77	54.63	1.12	1.37
	Y79AA1000268	43.67	51.54	45.57	1.18	1.04
50	Y79AA1000269	47.5	40.64	44.86	0.86	0.94
	Y79AA1000270	130.55	110.71	113.31	0.85	0.87
	Y79AA1000280	70.79	81.14	73.63	1.15	1.04
	Y79AA1000285	29.49	30.14	27.66	1	1
55	Y79AA1000295	35.22	30.73	33.72	1	1

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	VESEN1000137	54.08	51.03	54.97	0.94	1.02
	VESEN1000195	59.79	53.87	60.72	0.9	1.02
5	VESEN1000215	20.64	18.66	18.03	1	1
	VESEN1000279	182.2	194.3	161.47	1.07	0.89
	VESEN1000363	130.95	132.52	120.92	1.01	0.92
	VESEN1000388	47.19	34.28	39.83	0.85	0.85
	VESEN1000394	45.79	54.32	54.41	1.19	1.19
10	VESEN1000410	33.77	24.94	18.14	1	1
	VESEN1000411	40.62	43.06	55.86	1.06	1.38
	VESEN1000415	44.83	38.74	34.12	0.89	0.89
	VESEN1000440	77.99	76.99	81.63	0.99	1.05
15	VESEN1000452	55.97	59.97	56.08	1.07	1
	VESEN1000539	2520.3	1514.24	2475.51	0.6	0.98
	VESEN1000554	23.28	20.08	20.19	1	1
	VESEN1000557	65.42	99.26	91.48	1.52	1.4
20	VESEN1000575	33	42.85	36.23	1.07	1
	VESEN1000585	33.26	35.72	48.21	1	1.21
	VESEN1000592	14.7	10.84	12.9	1	1
	VESEN1000658	53.11	50.87	53.69	0.96	1.01
25	VESEN1000669	156.32	171.85	194.32	1.1	1.24
	VESEN1000743	30.01	26.82	23.72	1	1
	VESEN1000752	616.05	621.22	670.66	1.01	1.09
	VESEN1000761	102.22	128.97	144.62	1.26	1.41
	VESEN2000039	76.51	106.05	108.81	1.39	1.42
30	VESEN2000102	27.9	33.4	39.37	1	1
	VESEN2000164	75.95	85.45	57.77	1.13	0.76
	VESEN2000175	16.91	16.81	13.67	1	1
	VESEN2000186	47.71	41.92	47.45	0.88	0.99
35	VESEN2000199	210.67	251.42	274.58	1.19	1.3
	VESEN2000200	31.77	34.78	30.5	1	1
	VESEN2000204	42.87	27.57	46.4	0.93	1.08
	VESEN2000218	233.2	310.25	266.85	1.33	1.14
40	VESEN2000230	51.87	57.61	55.03	1.11	1.06
	VESEN2000272	66.18	71.44	47.62	1.08	0.72
	VESEN2000299	33.39	36.15	34.44	1	1
	VESEN2000323	122.34	153.67	120.03	1.26	0.98
45	VESEN2000327	25.64	38.53	30.18	1	1
	VESEN2000328	287.92	426.6	459.49	1.48	1.6
	VESEN2000330	210.17	635.42	400.1	3.02	1.9
	VESEN2000336	19.26	21	30.8	1	1
	VESEN2000354	29.98	44.87	45.25	1.12	1.13
50	VESEN2000378	214.41	200.95	233.25	0.94	1.09
	VESEN2000379	637.63	734.74	663.65	1.15	1.04
	VESEN2000397	33.94	34.23	25.21	1	1
	VESEN2000416	33.88	35.1	24.86	1	1
55	VESEN2000420	13.29	9.98	11.98	1	1

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	THYRO1001677	67.29	78.89	67.23	1.17	1
	THYRO1001683	134.37	158.27	162.12	1.18	1.21
	THYRO1001700	20.47	19.37	14.84	1	1
5	THYRO1001702	116	117.84	102.88	1.02	0.89
	THYRO1001703	48.06	45.1	43.24	0.94	0.9
	THYRO1001706	82.92	111.31	83.86	1.34	1.01
	THYRO1001721	34.49	36.81	37.64	1	1
10	THYRO1001725	156.38	135.31	116.05	0.87	0.74
	THYRO1001730	439.5	509.26	459.36	1.16	1.05
	THYRO1001738	45.7	45.43	46.39	0.99	1.02
	THYRO1001743	28.14	23.08	20.97	1	1
15	THYRO1001745	17.14	21.06	16.26	1	1
	THYRO1001746	31.83	30.99	31.95	1	1
	THYRO1001770	365.71	367.44	345.23	1	0.94
	THYRO1001772	123.83	151.39	150.86	1.22	1.22
20	THYRO1001778	125.3	136.46	114.03	1.09	0.91
	THYRO1001793	118.26	92.44	95.51	0.78	0.81
	THYRO1001796	35.4	35.03	31.41	1	1
	THYRO1001800	138.35	110.54	85.07	0.8	0.61
25	THYRO1001803	54.17	51.87	51.91	0.96	0.96
	THYRO1001809	68.41	58.62	37.24	0.86	0.58
	THYRO1001817	108.47	135.8	146.32	1.25	1.35
	THYRO1001819	52.11	82.26	80.67	1.58	1.55
30	THYRO1001828	4332.9	3758.25	2993.64	0.87	0.69
	THYRO1001854	310.97	325.29	244.91	1.05	0.79
	THYRO1001895	50.43	53.8	41.92	1.07	0.83
	THYRO1001907	95.74	95.16	76.26	0.99	0.8
35	TRACH1000006	28.12	33.72	27.35	1	1
	TRACH1000013	14.62	17.35	10.88	1	1
	TRACH1000074	91.95	104.52	91.83	1.14	1
	TRACH1000095	24.29	25.7	21.89	1	1
	TRACH1000102	89.04	112.46	84.84	1.26	0.95
40	TRACH1000108	18.93	22.62	9.17	1	1
	TRACH1000126	56.02	51.95	46.03	0.93	0.82
	TRACH1000146	55.68	61.38	44.09	1.1	0.79
	TRACH1000160	6.99	1.98	2.87	1	1
45	TRACH1000184	135.47	176.97	305.52	1.31	2.26
	VESEN1000004	30.77	39.29	26.61	1	1
	VESEN1000007	47.71	47.56	40.32	1	0.85
	VESEN1000013	92.2	108.84	158.66	1.18	1.72
50	VESEN1000028	152.94	139.07	153.59	0.91	1
	VESEN1000059	47.54	41	36.63	0.86	0.84
	VESEN1000100	43.33	51.75	59.44	1.19	1.37
	VESEN1000107	36.45	39.21	41.9	1	1.05
55	VESEN1000117	36.31	35.07	40.25	1	1.01
	VESEN1000122	47.58	33.92	57.09	0.84	1.2

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	THYRO1001426	295.65	347.22	278.53	1.17	0.94
	THYRO1001430	66.13	74.19	77.05	1.12	1.17
	THYRO1001434	32.32	68.71	43.27	1.72	1.08
5	THYRO1001456	36.49	32.57	41.27	1	1.03
	THYRO1001457	45.96	55.07	45.99	1.2	1
	THYRO1001458	37.02	45.23	50.19	1.13	1.25
	THYRO1001459	169.67	160.28	164.66	0.94	0.97
10	THYRO1001471	33.1	25.58	22.4	1	1
	THYRO1001478	23.72	21.89	21.41	1	1
	THYRO1001480	332.7	411.76	349.43	1.24	1.05
	THYRO1001481	120.8	159.14	137.5	1.32	1.14
15	THYRO1001487	120.62	158.11	135.86	1.31	1.13
	THYRO1001495	82.69	84.53	88.63	1.02	1.07
	THYRO1001498	24.31	46.88	45.76	1.17	1.14
	THYRO1001510	26.13	19.03	19.35	1	1
20	THYRO1001512	1869.6	2049.81	1317.06	1.1	0.7
	THYRO1001519	79.4	79.3	75.14	1	0.95
	THYRO1001522	89.46	101.52	59.9	1.13	0.67
	THYRO1001523	71.35	72.38	74.09	1.01	1.04
25	THYRO1001526	240.13	365.08	318.05	1.52	1.32
	THYRO1001529	90.07	80.02	57.67	0.89	0.64
	THYRO1001534	50.65	49.31	81.97	0.97	1.62
	THYRO1001537	166.69	165.39	144.11	0.99	0.86
	THYRO1001541	177.91	167.08	139.4	0.94	0.78
30	THYRO1001545	51.89	36.64	42.95	0.77	0.83
	THYRO1001559	145.05	146.22	132.42	1.01	0.91
	THYRO1001563	183.92	184.36	182.85	1	0.99
	THYRO1001570	30.95	31.51	31.27	1	1
35	THYRO1001573	65.95	39.85	42.27	0.61	0.64
	THYRO1001584	47.65	49.27	53.94	1.03	1.13
	THYRO1001593	82.04	72.74	68.9	0.89	0.84
	THYRO1001595	103.98	110.51	102.49	1.06	0.99
40	THYRO1001596	58.19	51.42	63.84	0.88	1.1
	THYRO1001602	87.72	85.88	72.86	0.98	0.83
	THYRO1001605	42.84	41.69	43.2	0.97	1.01
	THYRO1001608	121.51	57.35	96.56	0.47	0.79
45	THYRO1001617	117.84	136.24	110.76	1.16	0.94
	THYRO1001634	47.62	45.6	45.24	0.96	0.95
	THYRO1001637	208.17	214.42	182.7	1.03	0.88
	THYRO1001641	27.43	33.65	36.4	1	1
50	THYRO1001656	59.41	54.94	78.81	0.92	1.33
	THYRO1001658	164.89	276.29	264.03	1.68	1.6
	THYRO1001661	26.03	22.14	26.37	1	1
	THYRO1001671	40.07	38.84	37.89	1	1
	THYRO1001672	15.08	14.64	16.68	1	1
55	THYRO1001673	142.06	146.64	96.74	1.03	0.68

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	THYR01001104	49.54	71.39	81.87	1.44	1.65
	THYR01001109	26.9	24.31	17.57	1	1
5	THYR01001113	124.88	144.83	163.96	1.16	1.31
	THYR01001120	31.33	42.01	28.69	1.05	1
	THYR01001121	28.39	26.15	21.16	1	1
	THYR01001128	56.44	105.01	71.13	1.86	1.26
10	THYR01001133	129.29	141.09	109.23	1.09	0.84
	THYR01001134	6.12	48.8	3.2	1.22	1
	THYR01001142	13.49	9.24	6.46	1	1
	THYR01001173	69.76	76.75	76.64	1.1	1.1
	THYR01001175	18.21	19.94	20.86	1	1
15	THYR01001177	136.34	154.56	107.81	1.13	0.79
	THYR01001189	76.97	101.83	89.67	1.32	1.16
	THYR01001194	105.93	107.92	103.06	1.02	0.97
	THYR01001204	63.61	57.71	51.46	0.91	0.81
20	THYR01001205	351.28	452.06	359.35	1.29	1.02
	THYR01001213	77.39	103.68	84.02	1.34	1.09
	THYR01001224	136.69	152.12	149.46	1.11	1.09
	THYR01001237	40.47	50.92	42.87	1.26	1.06
25	THYR01001242	115.84	153.83	147.74	1.33	1.28
	THYR01001258	40.94	31.04	30.95	0.98	0.98
	THYR01001262	26.89	27.89	19.46	1	1
	THYR01001266	24.21	23.01	18.49	1	1
30	THYR01001271	33.6	39.06	47.64	1	1.19
	THYR01001287	618.09	675.7	490.75	1.09	0.79
	THYR01001290	8.16	5.43	5.68	1	1
	THYR01001291	59.43	91.42	60.21	1.54	1.01
35	THYR01001297	36.39	49.57	53.18	1.24	1.33
	THYR01001302	16.1	15.52	13.89	1	1
	THYR01001313	12.16	11.5	7.09	1	1
	THYR01001320	67.67	100.5	55.72	1.49	0.82
	THYR01001321	47.21	52.86	37.77	1.12	0.85
40	THYR01001322	24.97	27.56	20.65	1	1
	THYR01001327	22.74	18.51	20.74	1	1
	THYR01001336	67.89	117.68	67.41	1.73	0.99
	THYR01001347	30.07	15.91	22.89	1	1
45	THYR01001358	33.4	31.77	32.94	1	1
	THYR01001363	29.1	29.57	27.89	1	1
	THYR01001365	22.72	20.54	17.71	1	1
	THYR01001374	55.89	36.19	39.41	0.72	0.72
50	THYR01001401	67.18	66.1	79.9	0.98	1.19
	THYR01001403	36.96	35.79	36.33	1	1
	THYR01001405	60.25	66.49	51.3	1.1	0.85
	THYR01001406	99.99	118.14	125.66	1.18	1.26
55	THYR01001411	162.66	177.91	174.09	1.09	1.07
	THYR01001420	281.42	206.35	163.61	0.73	0.58

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	THYRO1000798	21. 24	23. 63	18. 2	1	1
	THYRO1000800	117. 04	315. 89	252. 61	2. 7	2. 16
	THYRO1000805	16. 57	11. 21	16. 09	1	1
5	THYRO1000815	65. 07	58. 64	52. 6	0. 9	0. 81
	THYRO1000829	58. 28	53. 39	49. 78	0. 92	0. 85
	THYRO1000835	27. 36	30. 54	24. 1	1	1
	THYRO1000843	62. 82	52. 11	57. 79	0. 83	0. 92
10	THYRO1000846	26. 19	21. 19	19. 35	1	1
	THYRO1000852	31. 05	29. 41	35. 84	1	1
	THYRO1000855	75. 51	92. 14	58. 07	1. 22	0. 77
	THYRO1000865	102. 88	116. 21	119. 95	1. 13	1. 17
15	THYRO1000866	45. 28	32. 65	45. 38	0. 88	1
	THYRO1000881	89. 25	87. 76	93. 28	0. 98	1. 05
	THYRO1000894	19. 5	18. 71	15. 61	1	1
	THYRO1000895	24. 84	14. 82	13. 39	1	1
20	THYRO1000916	86. 36	104. 38	80. 99	1. 21	0. 94
	THYRO1000917	1332. 1	1907. 82	1793. 1	1. 43	1. 35
	THYRO1000926	21. 25	19. 37	21. 25	1	1
	THYRO1000934	13. 71	17. 06	13. 29	1	1
25	THYRO1000951	18. 61	27. 82	26. 82	1	1
	THYRO1000952	28. 39	29. 36	28. 1	1	1
	THYRO1000956	15. 25	6. 93	8. 82	1	1
	THYRO1000960	12. 17	13. 58	16. 76	1	1
	THYRO1000961	37. 05	31. 39	30. 86	1	1
30	THYRO1000964	13. 96	17. 88	15. 56	1	1
	THYRO1000971	22. 43	35. 29	29. 46	1	1
	THYRO1000974	81. 82	86. 36	97. 53	1. 06	1. 19
	THYRO1000975	77. 2	76. 65	60. 02	0. 99	0. 78
35	THYRO1000983	57. 05	64. 85	57. 92	1. 14	1. 02
	THYRO1000984	51. 78	47. 85	48	0. 92	0. 93
	THYRO1000988	57. 88	51. 98	58. 3	0. 9	1. 01
	THYRO1000991	14. 79	9	10. 42	1	1
40	THYRO1000999	30. 62	26. 66	22. 67	1	1
	THYRO1001003	51. 62	45. 23	52. 45	0. 88	1. 02
	THYRO1001015	19. 88	10. 58	15. 95	1	1
	THYRO1001016	67. 86	67. 9	60. 96	1	0. 9
45	THYRO1001022	19. 18	19. 19	16	1	1
	THYRO1001031	157. 9	113. 8	73. 39	0. 72	0. 46
	THYRO1001033	20. 62	16. 08	14. 67	1	1
	THYRO1001062	38. 12	38. 16	24. 23	1	1
50	THYRO1001063	41. 75	35. 16	36. 06	0. 96	0. 96
	THYRO1001071	7. 28	6. 77	9. 2	1	1
	THYRO1001080	43. 21	35. 05	40. 67	0. 93	0. 94
	THYRO1001093	71. 48	81. 87	63. 33	1. 15	0. 89
55	THYRO1001100	20. 18	16. 25	13. 76	1	1
	THYRO1001102	36. 88	35. 13	38. 38	1	1

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	THYRO1000558	23.81	19.28	21.37	1	1
	THYRO1000569	74.05	68.57	73.64	0.93	0.99
	THYRO1000570	41.62	41.36	28.83	0.99	0.96
5	THYRO1000572	21.39	7.23	11.72	1	1
	THYRO1000573	19.98	17.86	16.98	1	1
	THYRO1000577	14.92	4.37	7.74	1	1
	THYRO1000580	35.2	44.75	27.1	1.12	1
10	THYRO1000584	51.21	39.64	46.27	0.78	0.9
	THYRO1000585	22.9	30.19	25.55	1	1
	THYRO1000596	13.06	12.04	12.16	1	1
	THYRO1000602	221.44	258.24	175.93	1.17	0.79
15	THYRO1000605	22.69	14.45	17	1	1
	THYRO1000615	20.64	20.24	22.7	1	1
	THYRO1000625	55.48	53.13	39.91	0.96	0.72
	THYRO1000636	67.42	74.2	69.95	1.1	1.04
20	THYRO1000637	28.99	27.15	20.45	1	1
	THYRO1000641	27.36	20.13	13.73	1	1
	THYRO1000657	25.52	29.79	32.29	1	1
	THYRO1000658	144.83	157.65	111.64	1.09	0.77
25	THYRO1000662	39.96	34.34	26.55	1	1
	THYRO1000666	35.48	38.48	23.93	1	1
	THYRO1000676	55.83	49.12	51.23	0.88	0.92
	THYRO1000678	17.96	23.23	20.9	1	1
	THYRO1000684	19.12	21.46	17.53	1	1
30	THYRO1000694	27.86	22.19	23.82	1	1
	THYRO1000699	89.92	90.86	76.51	1.01	0.85
	THYRO1000712	184.29	270.08	170.04	1.47	0.92
	THYRO1000715	143.06	353.21	194.25	2.47	1.36
35	THYRO1000716	33.56	44.54	28.99	1.11	1
	THYRO1000717	77.22	110.47	65.42	1.43	0.85
	THYRO1000723	11.31	19.35	9.86	1	1
	THYRO1000734	23.48	18.16	13.64	1	1
40	THYRO1000748	16.72	20.76	14.58	1	1
	THYRO1000755	58.84	52.18	63.38	0.89	1.08
	THYRO1000756	33.49	31	34.07	1	1
	THYRO1000776	50.16	42.39	50.04	0.85	1
45	THYRO1000777	23.87	35.27	19.82	1	1
	THYRO1000779	6.32	4.93	6.1	1	1
	THYRO1000782	117.27	125.31	142.84	1.07	1.22
	THYRO1000783	12.02	9.07	10.19	1	1
50	THYRO1000786	55.04	58.8	55.91	1.07	1.02
	THYRO1000787	19.96	12.46	13.7	1	1
	THYRO1000792	38.21	33.39	36.86	1	1
	THYRO1000793	17.33	13.84	12.2	1	1
	THYRO1000795	28.83	24.46	21.51	1	1
55	THYRO1000796	42.19	47.03	31.27	1.11	0.95

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	THYR01000187	77.72	82.92	67.1	1.07	0.86
	THYR01000190	75.91	65.3	68.76	0.86	0.91
5	THYR01000196	10.19	6.64	5.51	1	1
	THYR01000197	74.14	96.54	82.33	1.3	1.11
	THYR01000199	18.64	18.82	15.04	1	1
	THYR01000206	29.32	53.8	59.34	1.35	1.48
	THYR01000221	53.54	70.47	61.06	1.32	1.14
10	THYR01000222	38.81	43.06	43.08	1.08	1.08
	THYR01000228	45.93	36.69	32.8	0.87	0.87
	THYR01000241	97.96	115.66	91.36	1.18	0.93
	THYR01000242	39.55	36.22	38.32	1	1
15	THYR01000246	34.94	38.43	32.03	1	1
	THYR01000253	62.15	52.77	55.94	0.85	0.9
	THYR01000270	25.94	22.66	21.26	1	1
	THYR01000279	10.27	4.06	5.62	1	1
20	THYR01000285	91.85	96.96	103.11	1.06	1.12
	THYR01000288	44.89	35.11	51.9	0.89	1.16
	THYR01000296	51.1	53.97	49.14	1.06	0.96
	THYR01000320	52.48	54.06	54.2	1.03	1.03
25	THYR01000322	33.96	29.37	30.02	1	1
	THYR01000327	29.34	27.31	29.2	1	1
	THYR01000343	27.82	34.09	28.73	1	1
	THYR01000345	55.7	46.72	51.21	0.84	0.92
30	THYR01000358	9.41	7.7	6.57	1	1
	THYR01000368	19.77	17.83	17.93	1	1
	THYR01000375	67.03	68.23	60.2	1.02	0.9
	THYR01000381	15.46	15.89	24.65	1	1
35	THYR01000387	36.11	33.07	29.39	1	1
	THYR01000394	57.36	48.17	50.67	0.84	0.88
	THYR01000395	28.97	30.24	27.58	1	1
	THYR01000400	17.65	26.72	35.19	1	1
	THYR01000401	24.4	25.58	18.17	1	1
40	THYR01000407	16.4	16.1	17.58	1	1
	THYR01000420	46.85	54.92	41.53	1.17	0.89
	THYR01000438	50.01	46.6	46.81	0.93	0.94
	THYR01000452	42.1	54.28	49.65	1.29	1.18
45	THYR01000455	8.03	4.68	3.86	1	1
	THYR01000471	40.24	41.92	30.96	1.04	0.99
	THYR01000481	30.3	21.61	26.84	1	1
	THYR01000484	61.87	67.03	50.17	1.08	0.81
50	THYR01000488	20.11	25.07	21.14	1	1
	THYR01000501	22.62	16.34	20.31	1	1
	THYR01000502	16.19	23.43	22.05	1	1
	THYR01000505	13.77	12.09	10.63	1	1
55	THYR01000535	225.34	256.48	247.53	1.14	1.1
	THYR01000556	55.39	56.81	48.63	1.03	0.88

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	THYMU1000035	16.7	12.77	9.02	1	1
	THYMU1000037	19.37	14.84	13.58	1	1
	THYMU1000042	79.95	82.87	68.39	1.04	0.86
5	THYMU1000047	89.91	103.09	89.83	1.15	1
	THYMU1000080	40.1	18.75	23.34	1	1
	THYMU1000094	100.33	88.03	83.95	0.88	0.84
	THYMU1000109	3394.6	3204.08	2476.84	0.94	0.73
10	THYMU1000127	149.43	302.07	224.65	2.02	1.5
	THYMU1000130	50.99	54.49	39.21	1.07	0.78
	THYMU1000137	89.89	98.22	85.61	1.09	0.95
	THYMU1000146	41.95	48.71	54.46	1.16	1.3
15	THYMU1000159	75.18	54.17	55.6	0.72	0.74
	THYMU1000163	1438.2	1056.73	808.48	0.73	0.56
	THYMU1000167	37.01	32.25	20.73	1	1
	THYMU1000186	41.69	49.31	32.2	1.18	0.96
20	THYR01000017	64.39	60.8	44.03	0.94	0.68
	THYR01000026	34.47	79.34	36.05	1.98	1
	THYR01000034	27.76	31.8	20.76	1	1
	THYR01000035	35.4	40.29	62.06	1.01	1.55
25	THYR01000036	23.43	26.13	14.41	1	1
	THYR01000040	29.81	42.36	39.84	1.06	1
	THYR01000061	37.9	51.34	31.21	1.28	1
	THYR01000067	72.96	86.57	57.57	1.19	0.79
	THYR01000070	21.53	26.8	23.38	1	1
30	THYR01000072	36.97	46.63	32.18	1.17	1
	THYR01000084	44.96	56.29	56.67	1.25	1.26
	THYR01000085	49.35	66.17	67.44	1.34	1.37
	THYR01000086	11.1	15.74	10.31	1	1
35	THYR01000087	8.58	16.78	5.64	1	1
	THYR01000092	52.96	62.53	59.75	1.18	1.13
	THYR01000093	8.63	7.64	6.15	1	1
	THYR01000099	37.27	32.4	28.52	1	1
40	THYR01000107	24.41	21.8	26.8	1	1
	THYR01000111	21.9	17.22	25.47	1	1
	THYR01000121	20.79	17.72	18.35	1	1
	THYR01000124	8.23	6.71	9.71	1	1
45	THYR01000129	6.81	8.74	7.68	1	1
	THYR01000130	37.8	41.72	38.31	1.04	1
	THYR01000132	98.28	111.98	107.16	1.14	1.09
	THYR01000134	27.46	37.47	27.29	1	1
50	THYR01000144	28.27	22.09	27.89	1	1
	THYR01000155	12.36	10.14	12.44	1	1
	THYR01000156	51.42	49.18	46.2	0.96	0.9
	THYR01000163	65.44	73.1	64.89	1.12	0.99
55	THYR01000173	15.15	16	17.87	1	1
	THYR01000186	210.15	248.49	187.91	1.18	0.89

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	SKNMC1000014	92.84	113.71	75.96	1.22	0.82
	SKNMC1000018	54.44	47.21	45.13	0.87	0.83
5	SKNMC1000020	21.7	28.63	29.3	1	1
	SKNMC1000046	26.3	28.76	16.18	1	1
	SKNMC1000050	62.13	60.84	53.16	0.98	0.86
	SKNMC1000062	330.04	282.45	310.59	0.86	0.94
10	SKNMC1000075	17.55	17.79	15.1	1	1
	SKNMC1000082	22.09	43.54	26.48	1.09	1
	SKNMC1000091	55.45	72.12	67.23	1.3	1.21
	SKNMC1000099	15.48	12.31	17.8	1	1
	SKNMC1000104	19.72	17.77	17.75	1	1
15	SKNMC1000113	25.72	23.37	13.13	1	1
	SKNMC1000119	31.7	33.07	28.84	1	1
	SKNMC1000142	13.17	9.46	8.15	1	1
20	SKNMC1000170	18.9	35.22	13.42	1	1
	SKNMC1000178	36.62	37.44	34.52	1	1
	SKNMC1000194	30.86	16.71	25.89	1	1
	SKNMC1000198	34.11	45.61	47.99	1.14	1.2
	SKNMC1000225	26.81	26.95	26.7	1	1
25	SKNMC1000249	8.56	7.08	5.55	1	1
	SPLEN1000007	23.41	31.55	22.1	1	1
	SPLEN1000012	12.79	13.7	8.13	1	1
	SPLEN1000014	70.11	67.87	59.13	0.97	0.84
30	SPLEN1000036	594.83	558.47	477.44	0.94	0.8
	SPLEN1000059	18.59	12.25	11.97	1	1
	SPLEN1000068	70.13	64.3	54.94	0.92	0.78
	SPLEN1000072	32.05	40.98	29.41	1.02	1
35	SPLEN1000101	192.61	203.71	206.97	1.06	1.07
	SPLEN1000108	11.49	13.35	11.51	1	1
	SPLEN1000113	34.46	41.39	30.71	1.03	1
	SPLEN1000114	45.22	48.21	35.95	1.07	0.88
40	SPLEN1000132	34.95	36.95	42.28	1	1.06
	SPLEN1000135	343.33	311.82	277.27	0.91	0.81
	SPLEN1000136	66.82	82.96	80.88	1.24	1.21
	SPLEN1000141	76.01	69.18	98.16	0.91	1.29
	SPLEN1000164	30.6	24.91	22.66	1	1
45	SPLEN1000166	23.21	20.04	14.17	1	1
	SPLEN1000175	139.17	239.48	209	1.72	1.5
	SPLEN1000182	14.94	11.84	11.7	1	1
	SPLEN1000185	55.33	55.69	40.97	1.01	0.74
50	THYMU1000004	71.72	74.9	83.41	1.04	1.16
	THYMU1000009	94.13	77.8	90.28	0.83	0.96
	THYMU1000015	228.36	266.58	260.48	1.17	1.14
	THYMU1000016	85.88	95.7	65.55	1.11	0.76
55	THYMU1000023	20.69	15.9	18.99	1	1
	THYMU1000034	24.06	21.75	16.17	1	1

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	PLACE4000431	263.83	293	219.65	1.11	0.83
	PLACE4000443	20.1	8.3	13.07	1	1
	PLACE4000445	71.62	72.15	70.79	1.01	0.99
5	PLACE4000450	147.23	126.29	117.48	0.86	0.8
	PLACE4000455	62.39	48.46	44.08	0.78	0.71
	PLACE4000465	135.53	132.4	107.77	0.98	0.8
	PLACE4000466	1046.6	1140.39	799.29	1.09	0.76
10	PLACE4000472	121.8	126.78	113.94	1.04	0.94
	PLACE4000487	264.78	281.84	202.36	1.06	0.76
	PLACE4000489	48.45	36.77	38.35	0.83	0.83
	PLACE4000494	23.84	20.8	21.4	1	1
15	PLACE4000502	88.64	86.75	81.9	0.98	0.92
	PLACE4000521	323.04	404.56	285.14	1.25	0.88
	PLACE4000522	30.08	33.41	25.49	1	1
	PLACE4000537	29.74	23.24	26.85	1	1
20	PLACE4000548	42.55	40.31	87.2	0.95	2.05
	PLACE4000558	23.58	23.82	17.37	1	1
	PLACE4000581	61.42	69.56	63.92	1.13	1.04
	PLACE4000590	8.23	5	6.05	1	1
25	PLACE4000593	18.6	17.3	14.81	1	1
	PLACE4000612	41.95	53.37	32.82	1.27	0.95
	PLACE4000638	26.8	37.75	22.79	1	1
	PLACE4000650	27.3	16.4	33.1	1	1
30	PLACE4000651	36.82	39.3	38.22	1	1
	PLACE4000654	16.89	15.19	15.24	1	1
	PLACE4000670	16.92	13.19	15.71	1	1
	PLACE4000685	187.86	211.77	202.76	1.13	1.08
	PLACE4000687	16.21	22.59	13.56	1	1
35	PLACE5000003	21.87	28.54	21.39	1	1
	PLACE5000005	110.24	120.37	147.87	1.09	1.34
	PLACE5000019	13.52	8.92	10.26	1	1
	PLACE5000021	13.23	15.67	10.24	1	1
40	PLACE5000022	19.05	19.6	21.59	1	1
	PLACE5000024	34.25	21.2	21.25	1	1
	PLACE5000036	40.53	31.28	35.69	0.99	0.99
	PLACE5000059	352.37	319.61	357.62	0.91	1.01
45	PLACE5000076	47.21	58.74	32.77	1.24	0.85
	PLACE5000117	84.84	102.57	87.82	1.21	1.04
	PLACE5000143	23.5	22.78	23.28	1	1
	PLACE5000152	15.35	12.76	11.52	1	1
50	PLACE5000154	144.51	199.46	237.34	1.38	1.64
	PLACE5000155	89.36	81.41	79.91	0.91	0.89
	PLACE5000165	78.93	76.87	88.02	0.97	1.12
	SKNHC1000004	101.25	99.46	95.02	0.98	0.94
	SKNHC1000011	46.1	44.65	56.16	0.97	1.22
55	SKNHC1000013	16.47	25.9	14.18	1	1

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	PLACE4000106	42.56	54.08	43.9	1.27	1.03
	PLACE4000128	67.58	75.38	71.3	1.12	1.06
	PLACE4000129	29.93	24.81	23.87	1	1
5	PLACE4000131	782.86	711.57	587.61	0.91	0.75
	PLACE4000147	14.59	9.44	6.98	1	1
	PLACE4000156	163.18	267.14	159.69	1.64	0.98
	PLACE4000175	14.89	13.1	10.66	1	1
10	PLACE4000190	628.33	504.54	459.39	0.8	0.73
	PLACE4000192	28.59	35.85	25.67	1	1
	PLACE4000206	77.21	57.16	80.21	0.74	1.04
	PLACE4000211	79.65	101.47	63.07	1.27	0.79
15	PLACE4000214	26	17.61	16.37	1	1
	PLACE4000222	84.71	90.07	69.78	1.06	0.82
	PLACE4000223	22.62	14.09	14.21	1	1
	PLACE4000229	33.82	41.71	44.09	1.04	1.1
20	PLACE4000230	85.95	57.35	62.17	0.67	0.72
	PLACE4000233	73.47	68.6	63.73	0.93	0.87
	PLACE4000239	100.72	94.05	95.4	0.93	0.95
	PLACE4000247	34.55	28.2	23.03	1	1
25	PLACE4000250	51.35	59.3	45.61	1.15	0.89
	PLACE4000252	20.75	19.51	18.21	1	1
	PLACE4000259	250.69	228.39	179.09	0.91	0.71
	PLACE4000261	21.5	12.14	17.49	1	1
30	PLACE4000264	38.97	41.66	55.02	1.04	1.38
	PLACE4000269	47.1	37.8	32.16	0.85	0.85
	PLACE4000270	24.24	29.97	22.47	1	1
	PLACE4000281	211.22	274.25	267.89	1.3	1.27
35	PLACE4000300	24.03	26.73	16.79	1	1
	PLACE4000320	47.61	52.92	41.99	1.11	0.88
	PLACE4000323	70.64	77.63	73.66	1.1	1.04
	PLACE4000326	17.15	19.53	21.91	1	1
	PLACE4000344	20.15	17.55	19.07	1	1
40	PLACE4000347	42.11	35.66	37.07	0.95	0.95
	PLACE4000354	20.54	16.97	13.97	1	1
	PLACE4000367	19.08	21.62	13.54	1	1
	PLACE4000369	18.56	16.86	16.32	1	1
45	PLACE4000379	44.93	48.22	35.27	1.07	0.89
	PLACE4000387	28.48	18.41	23.23	1	1
	PLACE4000392	18.72	23.43	16.99	1	1
	PLACE4000399	417.14	435.88	349.26	1.04	0.84
50	PLACE4000401	13.12	19.41	12.77	1	1
	PLACE4000403	72.03	64.17	68.86	0.89	0.96
	PLACE4000411	59.36	51.92	41.47	0.87	0.7
	PLACE4000415	23.89	20.82	25.63	1	1
55	PLACE4000416	174.35	147.12	179.04	0.84	1.03
	PLACE4000424	19.71	16.25	17.79	1	1

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	PLACE3000309	48.91	70.41	44.97	1.44	0.92
	PLACE3000310	42.78	41.15	34.91	0.96	0.94
5	PLACE3000320	49.02	51.22	40.77	1.04	0.83
	PLACE3000322	3.16	0.57	1.79	1	1
	PLACE3000330	221.71	298.07	276.82	1.34	1.25
	PLACE3000331	138.49	160.45	123.87	1.16	0.89
	PLACE3000336	37.63	52.57	39.67	1.31	1
10	PLACE3000339	104.65	114.64	90.45	1.1	0.86
	PLACE3000341	114.82	141.48	110.97	1.23	0.97
	PLACE3000350	160.43	173.42	121.49	1.08	0.76
	PLACE3000352	47.6	57.89	40.6	1.22	0.85
15	PLACE3000353	61.18	67.39	56.69	1.1	0.93
	PLACE3000362	76	90.79	76.16	1.19	1
	PLACE3000363	16.4	27.93	15.48	1	1
	PLACE3000365	27.86	33.26	31.95	1	1
20	PLACE3000373	21.61	21.8	22.19	1	1
	PLACE3000374	31.32	41.25	22.3	1.03	1
	PLACE3000387	8.25	9.82	6.88	1	1
	PLACE3000388	64.69	77.9	44.31	1.2	0.68
25	PLACE3000399	112.65	133.33	109.36	1.18	0.97
	PLACE3000400	44.55	49.98	41.2	1.12	0.92
	PLACE3000401	490.44	600.16	510.14	1.22	1.04
	PLACE3000402	32.95	41.75	27.52	1.04	1
30	PLACE3000405	53.52	61.97	62.72	1.16	1.17
	PLACE3000406	35.93	38.71	42.23	1	1.06
	PLACE3000413	18.05	16.88	18	1	1
	PLACE3000416	29.64	28.6	31.16	1	1
35	PLACE3000425	50.28	54.42	57.61	1.08	1.15
	PLACE3000437	204.43	214.99	215.82	1.05	1.06
	PLACE3000455	70.16	96.18	109.86	1.37	1.57
	PLACE3000475	146.58	174.44	167.87	1.19	1.15
	PLACE3000477	71.98	48.83	46.22	0.68	0.64
40	PLACE4000003	23.25	25.51	19.95	1	1
	PLACE4000008	70.24	63.65	74.26	0.91	1.06
	PLACE4000009	43.12	37.54	47.82	0.93	1.11
	PLACE4000014	21.64	19.96	21.69	1	1
45	PLACE4000029	445.92	441.3	386.76	0.99	0.87
	PLACE4000034	90.28	104.98	136.62	1.16	1.51
	PLACE4000049	80.02	77.88	91.08	0.97	1.14
	PLACE4000052	28.33	17.89	22.57	1	1
50	PLACE4000062	33.61	37.15	30.93	1	1
	PLACE4000063	43.96	48.75	45.39	1.11	1.03
	PLACE4000089	59.97	63.79	58.77	1.06	0.98
	PLACE4000093	19.92	15.59	24.78	1	1
55	PLACE4000100	59.02	65.05	68.87	1.1	1.17
	PLACE4000103	23.19	22.13	16.6	1	1

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	PLACE3000052	367.89	489.29	371.66	1.33	1.01
	PLACE3000059	24.83	27.37	23.53	1	1
	PLACE3000067	59.74	70.09	61.55	1.17	1.03
5	PLACE3000069	93.54	91.2	73.59	0.97	0.79
	PLACE3000070	185.28	188.38	229.21	1.02	1.24
	PLACE3000103	65.17	61.33	54.93	0.94	0.84
	PLACE3000119	3.61	113.76	2.84	2.84	1
10	PLACE3000121	1345.4	1349.82	1036.69	1	0.77
	PLACE3000124	97.43	115.22	90.24	1.18	0.93
	PLACE3000135	11.35	10.57	7.29	1	1
	PLACE3000136	28.24	30.03	24.93	1	1
15	PLACE3000142	21.84	22.96	23.32	1	1
	PLACE3000145	738.49	611.83	451.51	0.83	0.61
	PLACE3000147	176.32	164.52	132.27	0.93	0.75
	PLACE3000148	4.21	2.12	1.47	1	1
20	PLACE3000154	35.29	60.89	56.38	1.52	1.41
	PLACE3000155	66.71	65.47	53.88	0.98	0.81
	PLACE3000156	18.22	36.57	17.58	1	1
	PLACE3000157	26.5	33.22	27.53	1	1
25	PLACE3000158	115.37	133.81	102.19	1.16	0.89
	PLACE3000160	136.75	99.83	114.37	0.73	0.84
	PLACE3000169	165.8	195.53	168.21	1.18	1.01
	PLACE3000181	127.79	141.83	120.13	1.11	0.94
	PLACE3000194	49.19	48.32	44.38	0.98	0.9
30	PLACE3000197	12.91	14.38	16.78	1	1
	PLACE3000199	14.71	10.56	10.36	1	1
	PLACE3000205	868.43	1057.62	753.31	1.22	0.87
	PLACE3000207	216.92	252.67	191.1	1.16	0.88
35	PLACE3000208	54.89	54.23	48.79	0.99	0.89
	PLACE3000213	33.26	40.23	30.49	1.01	1
	PLACE3000215	21.84	17.5	13.39	1	1
	PLACE3000218	13.41	11.66	9.09	1	1
40	PLACE3000220	114.32	105.77	107.52	0.93	0.94
	PLACE3000221	437.38	475.47	358.38	1.09	0.82
	PLACE3000225	48.79	61.71	52.94	1.26	1.09
	PLACE3000226	89.7	98.83	76.04	1.1	0.85
	PLACE3000230	26.28	30.38	53.41	1	1.34
45	PLACE3000231	34.3	53.63	46.02	1.34	1.15
	PLACE3000235	105.91	116.52	94.01	1.1	0.89
	PLACE3000242	62.07	67.67	62.09	1.09	1
	PLACE3000244	26.87	27.09	18.22	1	1
50	PLACE3000253	20.69	18.14	18.91	1	1
	PLACE3000254	59.24	71.24	61.14	1.2	1.03
	PLACE3000271	210.99	227.81	179.99	1.08	0.85
	PLACE3000276	22.03	19.83	15.86	1	1
55	PLACE3000304	283.54	350.52	301.89	1.24	1.06

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	PLACE2000302	45.55	58.9	46:59	1.29	1.02
	PLACE2000305	155.84	169.39	135.63	1.09	0.87
	PLACE2000317	32.82	20.28	20.47	1	1
5	PLACE2000324	41.35	38.7	36.15	0.97	0.97
	PLACE2000334	183.82	175.14	175.68	0.95	0.96
	PLACE2000335	136	196.92	135.48	1.45	1
	PLACE2000340	16.11	18.58	11.97	1	1
10	PLACE2000341	210.96	277.16	209.16	1.31	0.99
	PLACE2000342	166.99	114.91	107.63	0.69	0.64
	PLACE2000347	154.45	189.66	153.52	1.23	0.99
	PLACE2000357	79.73	96.17	102.29	1.21	1.28
15	PLACE2000358	256.57	305.62	211.22	1.19	0.82
	PLACE2000359	3.81	4.5	1.37	1	1
	PLACE2000366	99.02	132.98	84.55	1.34	0.85
	PLACE2000371	68.55	85.66	71.92	1.25	1.05
	PLACE2000373	43.55	47.22	46.75	1.08	1.07
20	PLACE2000374	23.53	32.86	28.16	1	1
	PLACE2000379	16.39	17.7	16.49	1	1
	PLACE2000386	1286.2	1518.95	1572.54	1.18	1.22
	PLACE2000388	50.87	49.22	36.25	0.97	0.79
25	PLACE2000392	319.51	348.31	353.39	1.09	1.11
	PLACE2000394	71.84	67.92	78.67	0.95	1.1
	PLACE2000398	30.22	38.93	35.07	1	1
	PLACE2000399	42.9	34.91	50.88	0.93	1.19
30	PLACE2000402	32.87	31.76	32.88	1	1
	PLACE2000404	58.57	67.45	73.86	1.15	1.26
	PLACE2000411	52.82	83.73	115.98	1.59	2.2
	PLACE2000418	50.46	65.86	56.08	1.31	1.11
35	PLACE2000419	102.69	137.4	124.2	1.34	1.21
	PLACE2000425	37.31	40.72	33.56	1.02	1
	PLACE2000427	21.66	25.88	27.14	1	1
	PLACE2000433	50.87	47.44	43.55	0.93	0.86
40	PLACE2000435	21.67	16.61	17.84	1	1
	PLACE2000438	19.41	25.68	29.91	1	1
	PLACE2000450	81.66	92.34	97.83	1.13	1.2
	PLACE2000455	18.01	22.78	20.07	1	1
	PLACE2000458	55.64	68.4	59.18	1.23	1.06
45	PLACE2000464	29.94	27.76	24.02	1	1
	PLACE2000465	213.81	251.88	207.06	1.18	0.97
	PLACE2000473	836.58	983.57	806.98	1.18	0.96
	PLACE2000477	16.45	29.51	16.05	1	1
50	PLACE3000004	87.82	83.69	80.62	0.95	0.92
	PLACE3000009	1606.2	1262.54	1148.63	0.79	0.72
	PLACE3000020	760.66	812.44	697.12	1.07	0.92
	PLACE3000029	217.19	259.74	201.46	1.2	0.93
55	PLACE3000038	41.85	54.63	40.3	1.31	0.96

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	PLACE2000022	46.15	49.28	51.9	1.07	1.12
	PLACE2000030	1694.2	2013.81	1419.12	1.19	0.84
	PLACE2000032	60.37	68.14	58.24	1.13	0.96
5	PLACE2000033	110.17	111.61	98.7	1.01	0.9
	PLACE2000034	57.78	58.66	54.29	1.02	0.94
	PLACE2000039	120.07	145.25	131.21	1.21	1.09
	PLACE2000043	346.07	354.6	252.31	1.02	0.73
10	PLACE2000044	23.97	29.58	29.62	1	1
	PLACE2000047	102.56	112.06	113.24	1.09	1.1
	PLACE2000050	71.89	90.85	58.04	1.26	0.81
	PLACE2000061	17.86	20.56	15.09	1	1
15	PLACE2000062	66.1	72.31	49.98	1.09	0.76
	PLACE2000072	17.99	18.55	16.31	1	1
	PLACE2000073	15.95	15.72	13.57	1	1
	PLACE2000097	213.87	252.44	257.53	1.18	1.2
20	PLACE2000100	78.87	88.68	57.86	1.12	0.73
	PLACE2000103	54.73	54.47	56.01	1	1.02
	PLACE2000106	135.55	158.65	131.58	1.17	0.97
	PLACE2000111	87.67	97.31	56.96	1.11	0.65
	PLACE2000115	17.7	16.8	12.26	1	1
25	PLACE2000118	108.86	115.04	163.24	1.06	1.5
	PLACE2000124	816.3	718.85	492.7	0.88	0.6
	PLACE2000132	13.87	16.39	13.31	1	1
	PLACE2000136	13.17	14.88	13.46	1	1
30	PLACE2000137	22.14	19.06	19.32	1	1
	PLACE2000140	98.61	148.4	126.8	1.5	1.29
	PLACE2000147	21.9	25.66	17.02	1	1
	PLACE2000153	14.7	11.12	14.66	1	1
35	PLACE2000164	20.49	18.37	20.21	1	1
	PLACE2000170	125.31	145.03	117.3	1.16	0.94
	PLACE2000172	17.54	16.31	16.59	1	1
	PLACE2000173	70.78	80.23	58.69	1.13	0.83
40	PLACE2000174	57.95	64.79	58.03	1.12	1
	PLACE2000176	43.43	41.92	40.16	0.97	0.92
	PLACE2000187	60.61	54.14	36.42	0.89	0.66
	PLACE2000216	59.22	72.23	56.49	1.22	0.95
	PLACE2000219	39.68	48.98	40.11	1.22	1
45	PLACE2000221	135.95	164.16	158.94	1.21	1.17
	PLACE2000223	5.6	6.49	3.36	1	1
	PLACE2000231	33.93	38.68	26.47	1	1
	PLACE2000235	98.28	112.79	92.76	1.15	0.94
50	PLACE2000246	94.02	98.46	83.22	1.05	0.89
	PLACE2000264	79.88	92.69	64.8	1.16	0.81
	PLACE2000274	35.59	37.8	39.08	1	1
	PLACE2000287	20.39	20.25	23.46	1	1
55	PLACE2000296	18.03	15.38	14.93	1	1

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	PLACE1011729	48.33	49.66	37.62	1.03	0.83
	PLACE1011741	42.11	42.21	26.14	1	0.95
	PLACE1011749	98.54	120.28	81.54	1.22	0.83
5	PLACE1011757	200.1	214.18	192.99	1.07	0.96
	PLACE1011762	31.14	34.16	27.67	1	1
	PLACE1011778	20.12	27.07	17.12	1	1
	PLACE1011783	79.37	103.33	80.61	1.3	1.02
10	PLACE1011795	62.52	74.53	49.61	1.19	0.79
	PLACE1011810	26.95	35.71	25.38	1	1
	PLACE1011824	94.06	261.39	163.66	2.78	1.74
	PLACE1011825	152.66	173.17	154.8	1.13	1.01
15	PLACE1011835	23.8	27.8	43.04	1	1.08
	PLACE1011836	120.62	135.76	141.71	1.13	1.17
	PLACE1011847	285.54	294.38	284.53	1.03	1
	PLACE1011855	22.68	27.11	28.04	1	1
20	PLACE1011858	36.79	39.34	31.86	1	1
	PLACE1011874	45.52	57.52	57.05	1.26	1.25
	PLACE1011875	21.91	19.56	19.38	1	1
	PLACE1011877	31.1	30.5	28.95	1	1
25	PLACE1011891	17.01	14.38	16	1	1
	PLACE1011896	9.03	4.47	5.22	1	1
	PLACE1011920	17.59	18.35	15.64	1	1
	PLACE1011922	14.43	22.14	26.94	1	1
30	PLACE1011923	318.75	346.7	273.3	1.09	0.86
	PLACE1011937	47.17	42.55	44.49	0.9	0.94
	PLACE1011939	72	86.06	70.82	1.2	0.98
	PLACE1011940	48.98	53.18	48.23	1.09	0.98
	PLACE1011962	47.89	49.93	54.67	1.04	1.14
35	PLACE1011964	25.14	30.6	30.08	1	1
	PLACE1011978	45.43	47.26	54.6	1.04	1.2
	PLACE1011980	45.74	54.67	59.79	1.2	1.31
	PLACE1011981	248.01	278.49	169.77	1.12	0.68
40	PLACE1011982	22.97	26.02	26.36	1	1
	PLACE1011995	53.57	49.73	50.05	0.93	0.93
	PLACE1012023	21.43	23.58	20.15	1	1
	PLACE1012026	22.39	20.44	21.38	1	1
45	PLACE1012031	29.21	36.05	24.28	1	1
	PLACE2000003	171.38	188.33	175.86	1.1	1.03
	PLACE2000005	17.08	13.58	14.3	1	1
	PLACE2000006	215.06	220.3	169.37	1.02	0.79
	PLACE2000007	32.91	35.43	30.39	1	1
50	PLACE2000011	77.03	82.84	64.01	1.08	0.83
	PLACE2000014	635.17	607.14	482.26	0.96	0.76
	PLACE2000015	37.29	41.56	31.59	1.04	1
	PLACE2000017	14.69	19.74	13.51	1	1
55	PLACE2000021	44.21	63.31	49.29	1.43	1.11

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	PLACE1011365	31. 47	37. 76	31. 3	1	1
	PLACE1011371	1393. 0	1511. 96	1103. 92	1. 09	0. 79
5	PLACE1011375	21. 17	21. 56	18. 08	1	1
	PLACE1011386	52. 19	45. 65	48. 93	0. 87	0. 94
	PLACE1011399	34. 74	30. 13	32. 85	1	1
	PLACE1011406	30. 27	36. 26	23. 82	1	1
	PLACE1011407	36. 4	42. 05	33. 38	1. 05	1
10	PLACE1011419	27. 2	44. 71	42. 78	1. 12	1. 07
	PLACE1011433	37. 98	29. 32	35. 67	1	1
	PLACE1011440	18. 75	14. 39	23. 74	1	1
	PLACE1011452	52. 41	64. 26	64. 32	1. 23	1. 23
15	PLACE1011465	25. 85	23. 7	25. 01	1	1
	PLACE1011472	25. 31	25. 72	28. 29	1	1
	PLACE1011477	589. 52	635. 4	447. 32	1. 08	0. 76
	PLACE1011478	115. 07	140. 27	106. 64	1. 22	0. 93
20	PLACE1011492	72	97. 27	57. 14	1. 35	0. 79
	PLACE1011498	17. 1	22. 26	16. 32	1	1
	PLACE1011501	17. 79	17. 26	18. 01	1	1
	PLACE1011503	9. 79	7. 51	7. 6	1	1
25	PLACE1011509	34. 49	28. 69	32. 22	1	1
	PLACE1011514	223. 19	247. 57	222. 01	1. 11	0. 99
	PLACE1011516	101. 6	143. 7	107. 63	1. 41	1. 06
	PLACE1011520	27. 41	24. 9	24. 32	1	1
	PLACE1011538	29. 14	23. 25	16. 78	1	1
30	PLACE1011555	23. 6	20. 53	25. 6	1	1
	PLACE1011561	24. 88	25. 71	37. 82	1	1
	PLACE1011563	37. 3	42. 35	32. 36	1. 06	1
	PLACE1011567	59. 12	55. 36	61. 57	0. 94	1. 04
35	PLACE1011569	110. 07	98. 25	92. 78	0. 89	0. 84
	PLACE1011576	107. 13	119. 56	82. 05	1. 12	0. 77
	PLACE1011586	58. 35	60. 15	53. 06	1. 03	0. 91
	PLACE1011635	126. 85	122. 41	79. 4	0. 96	0. 63
40	PLACE1011641	21. 95	15. 08	12	1	1
	PLACE1011642	96. 47	131. 07	98. 89	1. 36	1. 03
	PLACE1011643	27. 01	32. 97	23. 44	1	1
	PLACE1011646	311. 38	371. 3	292. 59	1. 19	0. 94
	PLACE1011649	39. 42	40. 05	38. 15	1	1
45	PLACE1011650	23. 59	25. 05	20. 03	1	1
	PLACE1011661	92. 23	95. 48	84. 22	1. 04	0. 91
	PLACE1011664	62. 94	63. 99	43. 95	1. 02	0. 7
	PLACE1011672	22. 5	26. 83	22. 49	1	1
50	PLACE1011675	20. 43	16. 82	14. 39	1	1
	PLACE1011682	34. 22	33. 4	46. 45	1	1. 16
	PLACE1011708	80. 24	92. 72	58. 81	1. 16	0. 73
	PLACE1011719	61. 08	68. 74	59. 57	1. 13	0. 98
55	PLACE1011725	39. 29	49. 72	30. 96	1. 24	1

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	PLACE1011041	31.57	35.69	21.03	1	1
	PLACE1011045	35.8	33.36	50.66	1	1.27
	PLACE1011046	34.53	36.4	27.94	1	1
5	PLACE1011054	87.07	91.74	90.14	1.05	1.04
	PLACE1011056	234.93	341.05	191.57	1.45	0.82
	PLACE1011057	50.32	61.94	47.85	1.23	0.95
	PLACE1011059	28.75	30.18	17.75	1	1
10	PLACE1011066	40.74	45.85	31.71	1.13	0.98
	PLACE1011087	126.9	97.7	103.08	0.77	0.81
	PLACE1011090	43.14	63.67	54.04	1.48	1.25
	PLACE1011109	72.28	109.31	75.78	1.51	1.05
15	PLACE1011114	40.38	55.61	44.13	1.38	1.09
	PLACE1011116	59.89	60.46	61.74	1.01	1.03
	PLACE1011122	23.85	22.65	27.32	1	1
	PLACE1011133	29.8	25.5	32.54	1	1
20	PLACE1011134	702.54	855.96	742.75	1.22	1.06
	PLACE1011143	18.05	26.08	15.76	1	1
	PLACE1011146	27.97	31.76	30.52	1	1
	PLACE1011160	27.65	29.84	36.02	1	1
25	PLACE1011165	29.12	25.53	32.58	1	1
	PLACE1011181	719.05	676.22	532.05	0.94	0.74
	PLACE1011185	128.61	184.54	128.11	1.43	1
	PLACE1011186	161.64	160.59	193.09	0.99	1.19
	PLACE1011203	26.77	26.81	25.39	1	1
30	PLACE1011214	745.48	767.22	632.47	1.03	0.85
	PLACE1011219	38.42	58.33	37.91	1.46	1
	PLACE1011221	31.09	40.42	44.12	1.01	1.1
	PLACE1011229	20.84	29.07	17.82	1	1
35	PLACE1011231	20.61	29.22	21.09	1	1
	PLACE1011236	553.05	663.66	472.59	1.2	0.85
	PLACE1011247	162.35	152.62	169.63	0.94	1.04
	PLACE1011263	28.78	26.07	29.73	1	1
40	PLACE1011273	22.76	23.17	22.03	1	1
	PLACE1011278	72.55	79.44	63.03	1.09	0.87
	PLACE1011289	23.79	33.63	31.25	1	1
	PLACE1011291	134.38	169.36	147.47	1.26	1.1
45	PLACE1011296	30.64	34.19	32.38	1	1
	PLACE1011310	29.41	24.49	28.94	1	1
	PLACE1011311	80.01	83.34	61.28	1.04	0.77
	PLACE1011321	63.79	59.32	62.43	0.93	0.98
	PLACE1011325	21.43	25.41	19.82	1	1
50	PLACE1011332	46.92	65.48	63.02	1.4	1.34
	PLACE1011340	96.73	105.33	93.22	1.09	0.96
	PLACE1011353	87.13	92.87	77.19	1.07	0.89
	PLACE1011360	73.75	62.56	56.98	0.85	0.77
55	PLACE1011364	40.61	42.21	47	1.04	1.16

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	PLACE1010716	26.71	33.85	35.18	1	1
	PLACE1010717	42.47	35.58	37.32	0.94	0.94
5	PLACE1010720	229.86	278.82	183.49	1.21	0.8
	PLACE1010739	32.44	33.25	22.11	1	1
	PLACE1010743	22.46	29.9	23.34	1	1
	PLACE1010752	18.27	19.55	21.76	1	1
10	PLACE1010761	104.86	122.36	130.93	1.17	1.25
	PLACE1010771	45.25	59.5	57.83	1.31	1.28
	PLACE1010784	20.56	18.65	15.58	1	1
	PLACE1010786	40.55	56.08	27.24	1.38	0.99
	PLACE1010789	27.09	24.19	22.66	1	1
15	PLACE1010800	34.72	39.52	36.8	1	1
	PLACE1010802	35.51	47.64	36.65	1.19	1
	PLACE1010811	21.55	29.77	27	1	1
	PLACE1010813	1287.7	1414.73	856.15	1.1	0.66
20	PLACE1010827	23	21.43	17.87	1	1
	PLACE1010833	33.54	35.98	30.25	1	1
	PLACE1010839	72	85.27	69.36	1.18	0.96
	PLACE1010856	86.05	87.34	90.48	1.01	1.05
25	PLACE1010857	57.56	61.88	60.74	1.08	1.06
	PLACE1010870	40.96	50.46	29.36	1.23	0.98
	PLACE1010877	46.38	39.36	49.51	0.86	1.07
	PLACE1010882	16.02	15.31	13.01	1	1
	PLACE1010891	16.64	15.08	17.9	1	1
30	PLACE1010896	30.16	23.93	23.2	1	1
	PLACE1010900	323.44	310.25	341.11	0.96	1.05
	PLACE1010916	20.94	23.68	16.14	1	1
	PLACE1010917	8.78	12.15	5.58	1	1
35	PLACE1010924	28.2	31.02	20.59	1	1
	PLACE1010925	59.4	63.62	79.68	1.07	1.34
	PLACE1010926	26.05	25.68	21.29	1	1
	PLACE1010942	73.09	63.9	54.96	0.87	0.75
40	PLACE1010943	114.11	144.64	114	1.27	1
	PLACE1010944	34.71	34.89	33.54	1	1
	PLACE1010947	50.97	56.41	46.75	1.11	0.92
	PLACE1010954	179.34	191.9	149.28	1.07	0.83
45	PLACE1010960	63.89	65.16	57.38	1.02	0.9
	PLACE1010965	63.35	54.39	57.39	0.86	0.91
	PLACE1010968	20	25	20.64	1	1
	PLACE1010978	52.43	69.4	55.7	1.32	1.06
50	PLACE1010982	40.37	51.27	37.65	1.27	0.99
	PLACE1010990	34.79	33.81	43.07	1	1.08
	PLACE1011017	114.63	113.45	90.55	0.99	0.79
	PLACE1011019	66.26	57.22	82.98	0.86	1.25
	PLACE1011026	61.83	68.6	53.2	1.11	0.86
55	PLACE1011032	44.7	54.48	36.43	1.22	0.89

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	PLACE1010341	14.28	17.1	15.38	1	1
	PLACE1010342	9.17	10.68	6.34	1	1
	PLACE1010346	25	30.67	21.84	1	1
5	PLACE1010362	35.78	48.63	28.59	1.22	1
	PLACE1010364	19.36	14.8	13.85	1	1
	PLACE1010368	32.4	36.5	44.4	1	1.11
	PLACE1010373	55.73	66.05	70.87	1.19	1.27
10	PLACE1010383	105.68	48.05	71.29	0.45	0.67
	PLACE1010385	6.58	5.18	2.96	1	1
	PLACE1010389	149.57	123.41	125.82	0.83	0.84
	PLACE1010401	15.79	15.26	16.26	1	1
15	PLACE1010410	28.15	37.81	34.22	1	1
	PLACE1010418	36.95	33.69	39.07	1	1
	PLACE1010425	16.45	18.69	17.64	1	1
	PLACE1010443	273.44	343.89	242.35	1.26	0.89
20	PLACE1010445	32.75	43.52	36.7	1.09	1
	PLACE1010481	19.87	17.5	20.62	1	1
	PLACE1010482	24.01	33.97	24.55	1	1
	PLACE1010491	35.66	27.01	27.99	1	1
25	PLACE1010492	97.73	111.42	92.48	1.14	0.95
	PLACE1010509	28.13	32.42	36.16	1	1
	PLACE1010518	98.12	123.6	114.12	1.26	1.16
	PLACE1010522	100.54	105.4	126.63	1.05	1.26
	PLACE1010529	32.36	59.45	68.71	1.49	1.72
30	PLACE1010547	21.89	34.68	16.06	1	1
	PLACE1010560	29.41	35.2	34.06	1	1
	PLACE1010562	23.25	30	23.72	1	1
	PLACE1010579	38.25	33.08	29.89	1	1
35	PLACE1010580	37.66	33.68	35.77	1	1
	PLACE1010599	92.73	121.5	90.34	1.31	0.97
	PLACE1010606	30.04	34.79	22.32	1	1
	PLACE1010616	46.86	69.11	81.69	1.47	1.74
40	PLACE1010622	28.24	38.42	25.03	1	1
	PLACE1010624	51.97	49.53	49.27	0.95	0.95
	PLACE1010628	40.93	49.51	35.73	1.21	0.98
	PLACE1010629	35.56	41.65	35.77	1.04	1
45	PLACE1010630	44.56	38.38	46.6	0.9	1.05
	PLACE1010631	27.36	20.75	17.51	1	1
	PLACE1010651	233.15	363.62	282.49	1.56	1.21
	PLACE1010661	22.85	27.37	31.66	1	1
	PLACE1010662	26.01	24.27	21.39	1	1
50	PLACE1010668	190.83	153.29	181.8	0.8	0.95
	PLACE1010702	59.8	61.23	55.06	1.02	0.92
	PLACE1010709	513.52	534.45	611.18	1.04	1.19
	PLACE1010713	60.05	51.29	72.19	0.85	1.2
55	PLACE1010714	19.13	16.92	16.72	1	1

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	PLACE1010039	19.95	22.68	18.66	1	1
	PLACE1010045	36.67	47.27	35.84	1.18	1
5	PLACE1010053	21.58	20.14	20.04	1	1
	PLACE1010060	38.71	56.37	32.78	1.41	1
	PLACE1010069	19.39	23.73	18.91	1	1
	PLACE1010070	20.01	20.72	21.19	1	1
10	PLACE1010074	165.08	128.25	150.68	0.78	0.91
	PLACE1010076	230.47	311.86	258.06	1.35	1.12
	PLACE1010078	31.79	36.25	37	1	1
	PLACE1010081	76.83	74.34	52.61	0.97	0.68
	PLACE1010083	17.09	21.13	24.34	1	1
15	PLACE1010089	20.95	24.75	17.3	1	1
	PLACE1010096	70.55	70.61	54.95	1	0.78
	PLACE1010102	39.64	55.68	62.77	1.39	1.57
	PLACE1010105	47.44	49.6	48.65	1.05	1.03
20	PLACE1010106	40.34	52.34	43.81	1.3	1.09
	PLACE1010130	40.29	44.66	30.55	1.11	0.99
	PLACE1010132	56.7	85.85	65.52	1.51	1.16
	PLACE1010134	20.51	15.99	20.46	1	1
25	PLACE1010139	415.41	498.57	421.38	1.2	1.01
	PLACE1010148	39.47	45.32	27.49	1.13	1
	PLACE1010152	29.32	28.44	29.8	1	1
	PLACE1010155	330.96	335.95	408.91	1.02	1.24
30	PLACE1010156	356.21	422.83	469.57	1.19	1.32
	PLACE1010161	37.57	44.88	36.68	1.12	1
	PLACE1010181	34.13	35.33	27.84	1	1
	PLACE1010194	24.35	22.48	25.49	1	1
	PLACE1010202	18.23	20.58	13.78	1	1
35	PLACE1010231	31.51	46.67	24.45	1.17	1
	PLACE1010235	45.66	38.3	32.66	0.88	0.88
	PLACE1010237	15.65	11.4	11.73	1	1
	PLACE1010251	45.81	45.52	23.23	0.99	0.87
40	PLACE1010261	32.21	38.4	27.38	1	1
	PLACE1010270	25.56	34.7	21.54	1	1
	PLACE1010273	20.85	22.99	16.74	1	1
	PLACE1010274	54.13	47.79	55.26	0.88	1.02
45	PLACE1010277	71.1	82.25	73.8	1.16	1.04
	PLACE1010293	66.72	98.76	53.8	1.48	0.81
	PLACE1010297	190.1	248.33	202.48	1.31	1.07
	PLACE1010300	45.22	39.59	27.21	0.88	0.88
50	PLACE1010310	424.98	529.37	453.64	1.25	1.07
	PLACE1010321	55.25	72.06	55.13	1.3	1
	PLACE1010324	25.61	25.41	17.98	1	1
	PLACE1010329	50.18	65.57	47.53	1.31	0.95
	PLACE1010330	55.1	76.67	55.02	1.39	1
55	PLACE1010335	77.42	150.11	159.9	1.94	2.07

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	PLACE1009665	36.25	41.13	28.7	1.03	1
	PLACE1009669	20.88	23.64	26.93	1	1
	PLACE1009670	27.37	30.25	30.27	1	1
5	PLACE1009708	24.07	37.18	35.56	1	1
	PLACE1009721	28.98	18.31	19.77	1	1
	PLACE1009731	35.52	28.93	25.46	1	1
	PLACE1009735	40.83	32.49	55.08	0.98	1.35
10	PLACE1009737	35.2	30.66	37.3	1	1
	PLACE1009741	18.21	26.46	34.29	1	1
	PLACE1009752	40.61	30.49	23.87	0.98	0.98
	PLACE1009763	49.08	51.16	54.68	1.04	1.11
15	PLACE1009766	41.6	36.03	39.03	0.96	0.96
	PLACE1009772	17.1	38.53	32.16	1	1
	PLACE1009782	44.04	32.84	31.11	0.91	0.91
	PLACE1009794	37.6	36.48	30.69	1	1
20	PLACE1009798	43.74	40.97	39.06	0.94	0.91
	PLACE1009845	17.16	29.15	19.05	1	1
	PLACE1009849	18.1	25.14	34.22	1	1
	PLACE1009857	24.47	21.19	22.73	1	1
25	PLACE1009861	157.04	147.3	184.61	0.94	1.18
	PLACE1009872	183.78	532.08	595.65	2.9	3.24
	PLACE1009877	88.74	127.12	82	1.43	0.92
	PLACE1009879	50.78	61.76	51.43	1.22	1.01
	PLACE1009886	26.47	30.9	26.46	1	1
30	PLACE1009888	40.97	27.49	37.88	0.98	0.98
	PLACE1009908	38.76	52.68	31.63	1.32	1
	PLACE1009919	72.43	157.7	91.06	2.18	1.26
	PLACE1009921	33.92	34.44	25.56	1	1
35	PLACE1009923	30.5	44.49	36.52	1.11	1
	PLACE1009924	99.41	54.1	55.42	0.54	0.56
	PLACE1009925	35.65	20.58	22.27	1	1
	PLACE1009931	90.33	98.46	76.54	1.09	0.85
40	PLACE1009935	25.18	24.01	15.51	1	1
	PLACE1009947	27.03	27.39	20.09	1	1
	PLACE1009961	24.51	25.25	21.83	1	1
	PLACE1009971	24.03	25.77	25.6	1	1
45	PLACE1009982	18.22	39.31	15.49	1	1
	PLACE1009992	9.19	11.11	8.64	1	1
	PLACE1009995	63.3	51.96	56.69	0.82	0.9
	PLACE1009997	43.45	50.16	41.93	1.15	0.97
	PLACE1010002	32.44	23.09	16.62	1	1
50	PLACE1010011	25.99	24.02	19.6	1	1
	PLACE1010013	15.52	18.47	13.26	1	1
	PLACE1010021	24.58	36.88	37.14	1	1
	PLACE1010023	43.52	60.71	61.14	1.39	1.4
55	PLACE1010031	52.61	51.96	38.23	0.99	0.76

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	PLACE1009388	49.9	74.96	45.01	1.5	0.9
	PLACE1009398	35.17	53.79	43.1	1.34	1.08
5	PLACE1009404	34.21	46.04	38.16	1.15	1
	PLACE1009410	26.75	35.99	28.94	1	1
	PLACE1009417	58.32	67.04	42.72	1.15	0.73
	PLACE1009424	34.63	42.13	42.37	1.05	1.06
10	PLACE1009434	24.85	24.58	24.73	1	1
	PLACE1009443	47.53	48.97	35.82	1.03	0.84
	PLACE1009444	31.89	50.08	34.61	1.25	1
	PLACE1009459	18.43	17.86	18.01	1	1
15	PLACE1009460	67.87	55.15	59.2	0.81	0.87
	PLACE1009468	61.44	73.54	56.01	1.2	0.91
	PLACE1009476	18.42	23.28	12.77	1	1
	PLACE1009477	38.48	38.41	40.05	1	1
	PLACE1009493	14.61	13.38	11.75	1	1
20	PLACE1009502	24.2	18.19	17.23	1	1
	PLACE1009524	18.39	17.32	15.6	1	1
	PLACE1009527	34.8	37.6	27.86	1	1
	PLACE1009531	202.28	205.54	217.56	1.02	1.08
25	PLACE1009535	41.74	48.48	38.26	1.16	0.96
	PLACE1009539	52.88	57.83	52.55	1.09	0.99
	PLACE1009540	74.03	82.77	96.82	1.12	1.31
	PLACE1009542	20.3	17.51	18.66	1	1
30	PLACE1009546	26.64	16.94	13.4	1	1
	PLACE1009556	19.56	22.58	16.96	1	1
	PLACE1009569	42.87	51.18	43.48	1.19	1.01
	PLACE1009571	28.33	26.69	25.34	1	1
35	PLACE1009573	55.23	54.49	56.76	0.99	1.03
	PLACE1009576	43.33	35.6	25.18	0.92	0.92
	PLACE1009580	66.98	54.33	52.85	0.81	0.79
	PLACE1009581	35.02	40.73	37.32	1.02	1
	PLACE1009587	21.2	22.71	17.17	1	1
40	PLACE1009593	69.23	95.74	64.2	1.38	0.93
	PLACE1009595	67.55	60.93	55.12	0.9	0.82
	PLACE1009596	31.43	27.75	24.05	1	1
	PLACE1009600	76.24	87.08	68	1.14	0.89
45	PLACE1009604	51.17	39.11	25.77	0.78	0.78
	PLACE1009607	105.68	136.82	84.46	1.29	0.8
	PLACE1009613	27.91	34.76	27.43	1	1
	PLACE1009621	70.93	83.59	74.98	1.18	1.06
50	PLACE1009622	51.58	101.79	73.07	1.97	1.42
	PLACE1009624	36.65	51.5	30.53	1.29	1
	PLACE1009637	14.3	19.67	15.18	1	1
	PLACE1009639	60.81	23.05	18.6	0.66	0.66
55	PLACE1009654	29.81	37.96	30.06	1	1
	PLACE1009659	49.17	66.83	38.29	1.36	0.81

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